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**Molecular Phylogenetic Studies in the Linaceae and *Linum*, with  
Implications for their Systematics and Historical Biogeography**

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**Molecular Phylogenetic Studies in the Linaceae and *Linum*, with  
Implications for their Systematics and Historical Biogeography**

**by**

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# **Molecular Phylogenetic Studies in the Linaceae and *Linum*, with Implications for their Systematics and Historical Biogeography**

Publication No. \_\_\_\_\_

Joshua Robert McDill, Ph.D.

The University of Texas at Austin, 2009

Supervisors: Beryl B. Simpson and Craig R. Linder

Best-known as the family of the cultivated flax, *Linum usitatissimum* L., the Linaceae is a small but ecologically diverse family of flowering plants, with approximately 250 species distributed throughout the temperate and tropical latitudes of the world. This work is an investigation of the systematics and biogeography of the family and a portion of its largest genus, *Linum*, using molecular phylogenetic methods. I collected DNA sequences of *rbcL* and *matK* genes from 51 species of Linaceae, representing all 14 genera, and combined them with data from 24 other families of the order Malpighiales in phylogenetic analyses. Results strongly support the monophyly of Linaceae and subfamily Linoideae in their current circumscriptions, but subfamily Hugonioideae is poorly supported. Molecular dating analysis suggests that the temperate Linoideae diversified in the Eocene or Oligocene, while tropical Hugonioideae diversified later, during the Miocene, perhaps ruling out Gondwanan vicariance as an explanation for their Panropical distribution. *Hugonia* and *Linum*, the largest genera in their respective subfamilies, are each found have multiple segregate genera nested within



them, indicating potential need for taxonomic revision of each subfamily. In Linoideae, I further investigate the phylogeny of a lineage that includes the yellow-flowered *Linum* sections *Cathartolinum*, *Linopsis*, and *Syllinum*, and the segregate genera *Cliococca*, *Hesperolinon*, *Radiola*, and *Sclerolinon*, to provide a framework in which to assess character evolution, classification, and biogeography. With data from four chloroplast markers (*matK*, *ndhF*, *trnK* 3' intron, *trnL-F* region) and the nuclear ITS, and extensive sampling from *Linum* section *Linopsis* from Eurasia, Africa, and the Americas, *Hesperolinon* and *Sclerolinon* are shown to be related to a lineage of Central American linums including *L. mexicanum* Kunth and *L. guatemalense* Benth., while *Cliococca* is affiliated with South American *Linum*. The phylogeny supports previous hypotheses of the evolution of some taxonomically important characters, and several well-supported lineages are identified which correspond to previously proposed taxonomic groupings. Results also provide evidence for a single trans-Atlantic disjunction and independent Old and New World colonizations of the southern hemisphere in yellow-flowered *Linum*, occurring during the Miocene.

## Table of Contents

List of Tables .....	xiii
List of Figures .....	xiv
Chapter 1: Introduction .....	1
Chapter 2: On the Relationship between Linaceae Subfamilies Linoideae and Hugonioideae .....	6
Abstract .....	6
Introduction .....	6
Materials and Methods .....	13
Results .....	18
Discussion .....	23
Conclusion .....	29
Chapter 3: Molecular Phylogenetics of of the Yellow-flowered flaxes, with Extended Sampling from <i>Linum</i> section <i>Linopsis</i> (Linaceae) .....	52
Abstract .....	52
Introduction .....	53
Materials and Methods .....	58
Results .....	63
Discussion .....	69
Conclusion .....	79
Chapter 4: On the Origins of Multiple Intercontinental Disjunctions in Yellow- Flowered <i>Linum</i> (Linaceae) .....	98
Abstract .....	98
Introduction .....	98
Materials and Methods .....	106
Results .....	114
Discussion .....	118
Conclusions .....	121

Appendix A: <i>rbcL</i> and <i>matK</i> alignment, Chapter 2.....	132
Appendix B: Voucher specimen information for Linaceae accessions utilized in Chapters 3 and 4.....	185
Appendix C: Chloroplast Data Alignment, Chapter 3 .....	195
Appendix D: Selected ITS Matrix, Chapter 3.....	278
References.....	299
Vita .....	309

## List of Tables

Table 2.1. The genera of Linaceae.....	31
Table 2.2. Historical treatments of Linaceae. ....	32
Table 2.3. Voucher specimens for taxa sampled for <i>matK</i> and <i>rbcL</i> .....	33
Table 2.4. Primer sequences. ....	36
Table 2.5. Sequence information. ....	37
Table 2.6. Estimated ages for nodes indicated in Fig. 2.13. ....	38
Table 3.1. Intraspecific classification of <i>Linum</i> sect. <i>Linopsis</i> . ....	81
Table 3.2. Primers used to amplify and sequence the <i>matK</i> gene from yellow- flowered <i>Linum</i> and outgroups. ....	83
Table 3.3: Data matrix characteristics. ....	84
Table 4.1: Data matrix characteristics for the reduced taxon set used for divergence time estimation and biogeographic hypothesis testing. ....	123
Table 4.2: SOWH test results for six constraints imposed on the reduced molecular dataset. ....	123
Table 4.3: Divergence times (mya) for lineages of yellow-flowered <i>Linum</i> , estimated by BEAST.....	124
Table A1: Species sampled for phylogenetic analyses in Chapter 3. ....	185

## List of Figures

Figure 2.1. Distributions of the eight genera of Linaceae subf. Linoideae.....	39
Figure 2.2. Distributions of the six genera of Linaceae subf. Hugonioideae.....	40
Figure 2.3. Strict consensus of 100, 158 MP trees (L=1725) based on rbcL.....	41
Figure 2.4. ML topology for rbcL recovered by GARLI.....	42
Figure 2.5. ML phylogram for rbcL.....	43
Figure 2.6. Strict consensus of 1898 MP trees (L=3691) based on matK.....	44
Figure 2.7. ML topology for matK recovered by GARLI.....	45
Figure 2.8. ML phylogram for matK.....	46
Figure 2.9. Strict consensus of 314 MP trees for combined data.....	47
Figure 2.10. ML topology for combined data recovered by RAxML using a partitioned model for combined tbcL and matK matrix.....	48
Figure 2.11. ML phylogram for combined data.....	49
Figure 2.12. Chronogram (maximum clade credibility topology) resulting from BEAST analysis with the split of Linaceae and Irvingiaceae set at 105 mya.....	50-51
Figure 3.1. Hypothesized evolutionary relationships among the various subgroups of <i>Linum</i> sect. <i>Linopsis</i> .....	85
Figure 3.2. Maximum likelihood topology for CPL matrix (-lnL=15848.70).....	86
Figure 3.3. 50% Majority-rule consensus of 30k post-burn-in trees sampled during Bayesian analysis of CPL matrix.....	87
Figure 3.4. Continuation of figure 3.3.....	88

Figure 3.5. ITS clone sequence ML bootstrap majority-rule consensus, part 1...	89
Figure 3.6. ITS clone sequence ML bootstrap majority-rule consensus, part 2...	90
Figure 3.7. ITS clone sequence ML bootstrap majority-rule consensus, part 3...	91
Figure 3.8. ITS clone sequence ML bootstrap majority-rule consensus, part 4...	92
Figure 3.9. ML bootstrap topology of selected ITS sequences.....	93
Figure 3.10. Majority-rule consensus from Bayesian analysis of ITS.....	94
Figure 3.11. ML topology for combined CPL+ITS matrix.....	95
Figure 3.12. Majority-rule consensus from Bayesian analysis of combined CPL+ITS data .....	96
Figure 3.13. Parsimony-optimized character mapping for four selected characters (topology=combined data ML).....	97
Figure 4.1. Distribution maps for yellow-flowered clade.....	125
Figure 4.2. Hypothetical historical biogeographic scenarios for yellow- flowered <i>Linum</i> .....	126
Figure 4.3. Parsimony mapping of geographic areas onto the ITS maximum likelihood topology. ....	127
Figure 4.4. Parsimony mapping of geographic areas onto the CPL maximum likelihood topology. ....	128
Figure 4.5. Parsimony mapping of biogeographic areas on combined (unreduced) CPL+ITS topology.....	129
Figure 4.6. Optimal ML topology for reduced data matrix, with parsimony mapping of geographic areas.....	130
Figure 4.7. Ultrametric majority-rule consensus of 27, 000 trees sampled during Bayesian analyses of yellow-flowered clade in BEAST.....	131

## Chapter 1: Introduction

The Linaceae is a small (ca. 250 species) but nearly cosmopolitan family of plants made famous by the cultivated flax, *Linum usitatissimum* L., which has been a commercially important source of fibers and seed oils since antiquity. Other flaxes, such as *L. perenne* L. and *L. grandiflorum* Desf., are cultivated as ornamentals, and a few (notably *L. album* Kotschy ex Boiss. and other species of *Linum* section *Syllinum* Griseb.) have been found to be sources of medicinally important lignans. In addition to *Linum* L., the family includes 13 additional genera, ranging from diminutive annuals of California's chaparral (*Hesperolinon* (Gray) Small, Sharsmith 1961) to canopy-forming trees in the humid, lowland forests of the Amazon basin (*Roucheria* Planch. and *Hebepetalum* Benth., Jardim 1999). They have been classified into two subfamilies (Dressler et al. in press): Linoideae (which includes *Linum* and the other annual and perennial genera of mostly temperate distribution) and Hugonioideae (tropical trees, shrubs, and lianas). In addition to their ecological and taxonomic diversity, the family displays a range of variation in several characters of biological and evolutionary interest, such as breeding system (both subfamilies include homostylous and heterostylous species), chromosome complement (with haploid numbers ranging from  $n=6$  to  $n=36$ ), pollen morphology (tricolpate to multiaperturate), floral pigmentation, and edaphic endemism. The Linaceae and *Linum* thus afford the opportunity to examine a number of evolutionary and biogeographic problems. In spite of the economic importance of the flaxes and a

history of detailed biosystematic study in the family, no previous attempt has been made to address the biogeography or evolution of Linaceae or *Linum* in a phylogenetic framework. Large-scale phylogenetic studies in Angiosperms (Savolainen 2000, Davis et al. 2005) have succeeded in placing Linaceae the order Malpighiales, but none have included broad sampling from the family.

The work presented here represents some preliminary contributions toward understanding evolutionary relationships in the Linaceae and *Linum*, using molecular phylogenetic analyses. In collaboration with Miriam Repplinger and Joachim Kadereit (both of Johannes Gutenberg University, Mainz, Germany), this led to an initial publication on the phylogeny of Linaceae subfamily Linoideae, with particular emphasis on the systematics, biogeography, and evolution of heterostyly in the five generally-recognized sections of *Linum* (McDill et. al. 2009). While several circumscriptions of *Linum* and its sections have been published in the last 150 years, the relationships among the sections and species of *Linum* and the other genera of Linoideae had not previously been assessed. The phylogeny of the subfamily was reconstructed using sequence data I collected from four chloroplast regions (*rbcL*, *trnK* 3' intron, *trnL-F* intron and intergenic spacer, and a small portion of the *ndhF* gene), in concert with additional *rbcL* and nuclear ITS sequences provided by Miriam Repplinger, with sampling representing all of the genera of Linoideae and the sections of *Linum*. Analyses revealed that *Linum* is divided into two major lineages, designated the Blue-Flowered and Yellow-Flowered clades. The two largest sections of *Linum*, sects. *Linum* and *Linopsis*, were found to be paraphyletic, with the remaining sections nested within them. *Linum* itself was also shown to be non-



monophyletic, the segregate genera *Cliococca*, *Hesperolinon*, *Radiola*, and *Sclerolinon* being interspersed among the yellow-flowered species of *Linum*. Results from that study also implied that Eurasia was the ancestral area in which the lineages of *Linum* initially diversified.

Another finding of those early analyses was that the tropical subfamily, Hugonioideae, was suggested to be paraphyletic, but this conclusion was tempered by the fact that very few Hugonioid taxa were sampled, with only three of the subfamily's six genera represented with data from a single marker, *rbcL*. This led to the analyses presented here in Chapter 2, an examination of the relationship between Linaceae subfamilies Linoideae and Hugonioideae, with data from two chloroplast markers, *rbcL* and *matK*, and sampling from 51 species representing all 14 genera of the family. These data were combined with published, publicly available sequences from representatives of 24 other families of Malpighiales, so that the monophyly of Linaceae could be assessed. Results of phylogenetic analyses using maximum parsimony, maximum likelihood, and Bayesian methodologies indicate that the two subfamilies may be monophyletic sister lineages, although Hugonioid monophyly might still be in question.

Chapter three revisits the phylogeny of *Linum* with an emphasis on *Linum* section *Linopsis*. Section *Linopsis* is the largest and most widespread of *Linum*'s five traditional sections — while the other four sections of *Linum* are essentially confined to Eurasia, *Linopsis* has diversified extensively in Africa and the Americas. *Linopsis* is also the most thoroughly-studied section taxonomically, thanks to the work of C.M. Rogers and his student R. Mildner, who produced revisionary studies of African and American flaxes

(Rogers 1963, 1981a, 1981b, 1984; Rogers and Mildner 1976) as well as a systematic classification for the entire section (Rogers 1982). The previous analyses published by McDill et al. (2009) indicated that the members of section *Linopsis* were part of a loosely-characterized “yellow-flowered” lineage that was sister to *Radiola linoides* Roth, and included *Linum* sections *Cathartolinum* Griseb. and *Syllinum* as well as the segregate genera *Cliococca* Bab., *Hesperolinon*, and *Sclerolinon* C.M. Rogers. The phylogeny of this lineage is reconstructed using maximum likelihood and Bayesian methods, with data from four chloroplast markers and the nuclear ITS region and extensive taxon sampling from *Linum* section *Linopsis* from Europe, Africa, and the Americas. The goals of this study were to clarify the relationships of the segregate genera, assess the classification and character evolution hypotheses that have been proposed for section *Linopsis*, and provide a phylogenetic framework for the biogeographic analyses presented in Chapter 5.

In Chapter 4, the phylogeny of the yellow-flowered clade is used to identify the intercontinental disjunctions that have taken place during their diversification, and to test different historical biogeographic hypotheses for the origins of these disjunctions based on their phylogenetic predictions and divergence time requirements. Because the data and analyses presented in Chapter 3 could not conclusively resolve the relationships among the species groups of South American, North American, and South African *Linum* (including the segregate genera), a parametric bootstrapping method was used to test different hypotheses of relationships aimed at determining the number of trans-Atlantic and trans-tropical disjunctions necessary to explain the current distribution of this lineage. A Bayesian method of divergence time estimation, employing calibration points

based on previous estimates for the age of the family and a *Linum* pollen fossil from the late Eocene, is used to determine whether the timing of the apparent disjunctions is compatible with historical biogeographic scenarios involving vicariance, land bridge migration, or recent long-distance dispersals.

The following three chapters of this dissertation will ultimately be published as independent papers in scientific journals. Each has been formatted with its own abstract, introduction, materials and methods, results, tables, figures, and discussion. A single list of references has been assembled for all of the chapters combined.

## **Chapter 2: On the Relationship between Linaceae Subfamilies Linoideae and Hugonioideae**

### **ABSTRACT**

The relationship between Linaceae subfamilies Linoideae (temperate) and Hugonioideae (tropical) are examined with data from two chloroplast regions and sampling representing all 14 commonly-recognized genera of Linaceae as well as representatives of 24 families of the order Malpighiales. The Linaceae and both of its subfamilies are found to be monophyletic, although the monophyly of Hugonioideae is not well supported. The families of Cronquist's order Linales – Ctenolophonaceae, Erythroxylaceae, Humiriaceae, and Ixonanthaceae – are not found to share recent common ancestry with Linaceae, though relationships of families are not well resolved in our analyses and no definitive sister-group to Linaceae can be identified. Both *Hugonia* and *Linum*, the largest genera of their respective subfamilies, are found to be non-monophyletic, each with several segregate genera nested within them. Average divergence time estimates indicate that the extant Linaceae began to diversify in the Paleocene, with the majority of species originating in the Miocene or later.

### **INTRODUCTION**

The Linaceae are a relatively small but subcosmopolitan family of flowering plants. Including the cultivated flax (*Linum usitatissimum* L., the source of linen fiber and linseed oil), the family comprises approximately 270 species, divided into 14 genera in

two subfamilies: the Linoideae and the Hugonioideae (Dressler et al., in press). As a whole, the Linaceae is only weakly diagnosable by a suite of morphological characters, none unique among angiosperms: simple, entire leaves that vary among species in their arrangement but are most commonly alternate, flowers that are hermaphroditic, actinomorphic, and pentamerous (four-merous in two species), with caducous petals often marked by pigmented lines or bands, five or ten stamens that are fused basally into a cup or tube that surrounds the ovary, and a syncarpous superior ovary of five carpels (but the number of carpels is reduced in many taxa).

With approximately 210 species, Linoideae is the larger subfamily, and encompasses the type genus, *Linum* L., and seven other genera (Table 2.1). Their distributions are primarily temperate, as shown in Fig. 2.1, although the ranges of some species do extend into, or are entirely within, tropical latitudes. Those species of Linoideae that inhabit the tropics tend to occur in relatively open, arid habitats and highlands, not the canopied, lowland mesic forests more typically considered as tropical habitats. Linoideae are herbaceous annuals or shrubby perennials, and are differentiated from Hugonioideae by the number of fertile stamens (five in Linoideae, alternating with structures interpreted as staminodia in some species) and by their dehiscent, capsular fruits. In the first detailed global treatments of Linaceae, Planchon (1847, 1848) recognized four of the genera now placed in subfamily Linoideae: *Anisadenia* Wall., *Linum*, *Radiola* Hill., and *Reinwardtia* Dumort. *Tirpitzia* Hallier f. was segregated from *Reinwardtia* by Hallier (1921) to accommodate a species with long corolla tubes not seen in other *Reinwardtia*. The remaining three genera were segregated from *Linum* in

recognition of their morphological distinctiveness: *Cliococca* Bab. (Babington 1842, Rogers and Mildner 1971), with its unique, closely-packed, needle-like leaves and indehiscent fruit, *Hesperolinon* (A. Gray) Small with its reduction in the number of carpels from five to three or two (Small 1907, Sharsmith 1961), and *Sclerolinon* C.M. Rogers (Rogers 1966) with its bicarpellate fruits that break apart into four indehiscent nutlets (as opposed to the dehiscent capsules found in other Linoideae).

The Hugonioideae, with approximately 57 species in six genera, are mostly confined to the tropics (Fig 2.2). They are trees, shrubs, and lianas that tend to inhabit moist lowland forests, either as part of the understory or the canopy (Van Hooren and Nootboom 1984, Jardim 1999). Several species occur in arid habitats, such as *Hugonia orientalis* Engl., a species of the Sandveld vegetation in southern Africa (Schmidt et al. 2007). In addition to their growth habit and tropical distribution, Hugonioideae are characterized as having ten fertile stamens (in two ranks of different height). The fruits are fleshy and are often described as drupes (pyrenes in *Durandea* and some *Hugonia*), although they are derived from multiple carpels and usually contain more than one seed (van Hooren and Nootboom 1984, Jardim 1999).

The generic taxonomy of Hugonioideae proceeded with the early recognition of *Hugonia*, followed by removal of variant species to small, segregate genera. *Hugonia* L. was known to Linnaeus as an Old World genus of trees and woody vines. Planchon recognized *Roucheria* Planch. based on collections from South America (two species, *R. humiriifolia* Planch. and *R. calophylla* Planch.) and India (*R. griffithiana* Planch.) in which the number of carpels was reduced from that observed in *Hugonia* (Planchon

1848). At the same time, Planchon segregated the genus *Durandea* Planch., which has pyrenaceous fruits, smaller flowers, and lacks the large, pinnatifid stipules found in *Hugonia* (Planchon 1848, Winkler 1931, Jardim 1999). *Durandea* was later treated as a subgenus of *Hugonia* (Baillon 1878, Van Hooren and Nooteboom 1984). In 1862, Planchon's *Roucheria humiriifolia* served as the basis for another new genus, *Hebepetalum* Benth., due in part to its distinctive leaf venation (Bentham and Hooker 1862) and internal corolla pubescence; subsequent shuffling of taxa between *Hebepetalum* and *Roucheria*, and the description of new species of each, are reviewed in detail by Jardim (1999). Hallier segregated *Philbornea* Hallier f. from *Durandea*, based in part on its inflorescence structure and reduced number of carpels from five to three or four (Hallier 1912). Hallier later transferred *Roucheria griffithiana* to another new genus, *Indorouchera* Hallier f., noting that it is a tendrillate vine of the Nicobar islands and Malaysia, whereas *Roucheria* species are South American trees that lack tendrils (Hallier 1921, Jardim 1999).

Several additional woody tropical genera have been treated as members of Linaceae due to perceived similarities to members of Hugonioideae. Table 2.2 summarizes four historical classifications schemes for Linaceae along with the current classification (Dressler et al. in press). The expanded circumscriptions of Bentham and Hooker (1862), Baillon (1878), and Winkler (1931) included genera alternatively placed in the Erythroxylaceae, Ctenolophonaceae, Ixonanthaceae, and Humiriaceae. Later treatments usually excluded these groups from Linaceae, based on studies of floral morphology and wood and leaf anatomy (Van Hooren and Nooteboom 1984), but still

usually considered them close relatives to Linaceae. Cronquist recognized the perceived similarities of these families to Linaceae by classifying them all in his order Linales (Cronquist 1988).

Investigations of angiosperm phylogeny have provided some additional evidence for the exclusion of many tropical components from Linaceae and have clarified the ordinal placement of the family, in addition to supporting the close relationship of Hugonioideae and Linoideae. All of the families of Cronquist's Linales are now placed in the large order Malpighiales (APG II 2003). Analyses of sequence variation in the chloroplast genes *rbcL* and *atpB*, and the nuclear ribosomal gene 18S, found that the Humiriaceae (represented by *Humiria* and *Vantanea*), Ixonanthaceae (*Ochthocosmus* and *Ixonanthes*), and Erythroxylaceae (*Erythroxylon* and *Nectaropetalum*) were dispersed throughout the Malpighiales, and do not appear to share recent common ancestry with Linaceae (Savolainen et al. 2000). A more strictly defined Linaceae, represented by one sample each from *Hugonia*, *Reinwardtia*, and *Linum*, was shown to be monophyletic in that study. The closest relative of Linaceae remains unknown; clades that have appeared as sister groups to Linaceae in various analyses include Violaceae + Passifloraceae + Turneraceae (Savolainen et al. 2000), Picrodendraceae (Chase et al. 2002), Balanopaceae + Trigoniaceae + Dichapetalaceae + Euphroniaceae + Chrysobalanaceae (Davis and Chase 2004), or Irvingiaceae (Davis et al. 2005). Sampling of Linaceae in these studies has been very limited, with one *Reinwardtia* and one or two species of *Linum* representing Linoideae, and one species of either *Hugonia* or *Durandea* representing Hugonioideae. The sister relationship of Linoideae and Hugonioideae has been found



consistently in phylogenetic studies that included members of both subfamilies.

However, many of the genera of Linaceae (whether currently accepted or historically supposed) have not been sampled together in molecular phylogenetic analyses, leaving our understanding of the membership of Linaceae and the relationships among its genera incomplete.

The Hugonioideae (or Hugoniaceae) and Linoideae (Linaceae s.s.) could be considered an example of an angiosperm “family pair” in which one member is mainly tropical and the other is temperate to cosmopolitan in distribution (Judd et al. 1994). In several such family pairs, such as the Apocynaceae/Asclepiadaceae, it has been shown that the first, mainly tropical member, is paraphyletic, with the second, more temperate family derived from within it (Potgeiter and Albert 2001). In other pairs, such as the Clusiaceae/Hypericaceae and Araliaceae/Apiaceae, more complex patterns of relationship have been revealed, implying that the temperate and tropical groups are each monophyletic yet derived from a common ancestry (Gustafsson et al. 2002, Plunkett et al. 1997). Results from phylogenetic studies of such family pairs can have important consequences for family circumscriptions and recognition (if monophyly is considered an necessary qualification for family recognition). Though Hugoniaceae is occasionally recognized as a distinct family, and numerous morphological and anatomical differences exist that distinguish it, its relationship to Linaceae s.s. (= Linaceae subf. Linoideae) has not been specifically investigated in any phylogenetic analyses.

A preliminary view of Linaceae phylogeny with *rbcL* sequences representing all genera of Linoideae plus several Hugonioid taxa (three species of *Hugonia*, one

*Durandea*, and two *Roucheria*) was presented by McDill et al. (2009). An interesting finding of their analysis was that the Hugonioideae appeared paraphyletic, with a well-supported, monophyletic Linoideae nested within it. This result could have important implications for the circumscription of Linaceae, supporting the idea that Hugoniaceae should not be recognized as a distinct family. However, it could also represent an anomalous result stemming from lack of phylogenetic signal, inadequate taxon sampling from Hugonioideae, or inadequate outgroup selection leading to mis-rooting of the Linoideae/Hugonioideae clade. McDill et al. (2009) went on to provide a detailed account of relationships in Linoideae, with additional data and taxon sampling from a broad sample of that subfamily, but generic relationships in Hugonioideae, and its disposition relative to Linoideae, were not explored further.

In the present study we revisit the question of the relationship between Linoideae and Hugonioideae. Taxonomic sampling from Hugonioideae has been increased to represent all of the genera currently placed in the subfamily, and a somewhat larger fraction of its largest genus, *Hugonia*. To complement the *rbcL* data, a parallel matrix of sequences of the chloroplast gene *matK* has been added. Outgroup taxa were chosen to represent most of the other families of Malpighiales (as sampled by Tokuoka and Tobe 2006), including the Ctenolophonaceae, Erythroxylaceae, Humiriaceae, and Ixonanthaceae, which have been hypothesized to be closely related to (or members of) the Linaceae in the past (i.e., as members of Winkler's 1931 expanded Linaceae, or Cronquist's order Linales). The questions this study seeks to address are summarized as follows:

1. Is the Linaceae monophyletic in its current circumscription (that of Dressler et al., in press)?
2. Do any other members of Cronquist's order Linales or the "expanded" Linaceae of Bentham and Hooker, Baillon, or Winkler share recent common ancestry with Linaceae, to the exclusion of other members of Malpighiales?
3. Is the Hugonioideae monophyletic and potentially recognizable as a family distinct from Linaceae?
4. What are the generic relationships in the Hugonioideae?

## **MATERIALS AND METHODS**

**Taxon and Marker Selection** — Fifty-one species were chosen to represent the 14 currently recognized genera of Linaceae. Each genus is represented by multiple species when possible, and the largest genera (*Linum* and *Hugonia*) were sampled to approximate the breadth of their taxonomic diversity and geographic range. Sequences of the chloroplast genes *matK* and *rbcL* were obtained from these selected taxa (some *rbcL* having been sequenced previously by McDill et al. [2009]), and combined with sequences representing 24 families of the order Malpighiales, available from GenBank. These markers were selected due to their general usefulness in elucidating angiosperm phylogeny (Savolainen et al. 2000, Hilu et al. 2003) and for the availability of sequences from a broad sample of taxa from the Malpighiales. Outgroups for tree rooting purposes were selected from the six orders that, along with Malpighiales, comprise the Eurosoid I

lineage (APG 2003). Taxon sampling from Malpighiales and outgroup selection represent a subset of taxa sampled by Tokuoka and Tobe (2006). Table 2.3 provides a complete list of taxa sampled, including GenBank accession numbers for new and previously-published sequences.

**DNA Extraction, PCR, and Sequencing**—DNA was extracted from fresh leaves or herbarium specimens using a modified Doyle and Doyle CTAB protocol (Loockerman and Jansen 1996). Two to six microliters of a 1:10 dilution (in TE) of each extract were used as template in PCR reactions. The primers used for amplification and sequencing of *rbcL* and *matK* that are new to this study are provided in Table 2.4, along with previously published primer sequences. PCR was executed in 25 $\mu$ L reaction volumes, the master mix including (per reaction): 2.5 $\mu$ L 10X Triton-X reaction buffer, 200  $\mu$ M dNTPs, 10pM each primer, 2-3  $\mu$ L 25 mM MgCl<sub>2</sub>, and *Taq* polymerase. Addition of BSA (3 $\mu$ L of 3.3% w/v BSA, Savolainen et al. 2000) to the reaction mix enhanced amplification from some samples. The thermal cycling program for amplification consisted of the following steps: initial denaturation at 95°C for 4 min, followed by 34 cycles with denaturation at 95°C for 30 sec, annealing at 50°C for 30 sec, and extension at 72°C for 30 sec. Cycles were followed by a final extension step at 72°C for 6 min. PCR products were verified by electrophoresis in 1.5% agarose gels, and cleaned using either QiaQuick columns (Qiagen Inc.) or Centri-Sep columns packed with G-50 Sephadex (Princeton Separation, Inc.), according to the manufacturer's instructions, or using the EXO-SAP protocol based on Werle et al. (1994).

Amplified fragments were sequenced with forward and reverse dye terminator reactions using the original amplification primers or internal primers as necessary for each marker and BigDye 3.1 chemistry (Applied Biosystems). Sequencing was done at The University of Texas at Austin on either an MJ Research BaseStation sequencer (MJ Research, Waltham, Massachusetts, discontinued) operated by the author, or an ABI 3730 DNA Analyzer at the Institute for Cell and Molecular Biology Core Facility.

All forward and reverse sequence reads were assembled and edited in Sequencher 4.1.2 or 4.5 (GeneCodes Corp. 2005). Alignment of *rbcL* was achieved manually in MacClade 4.08 (Maddison and Maddison 2005), while *matK* sequences were aligned using the MAFFT tool (Kato et al. 2002) to align sequences using their amino acid translation, followed by adjustment in MacClade. When data from either marker could not be obtained from a specific taxon, that taxon was included in the combined analysis with the missing sequence coded as missing data.

**Phylogenetic Analyses** — Analyses using maximum parsimony (MP) as the optimality criterion were conducted with PAUP\* 4.0b10 (Swofford 2002). The heuristic search algorithm was engaged with 500 random addition replicates and tree bisection-reconnection (TBR) branch-swapping, saving and swapping to completion on all minimal trees found. Bootstrap analyses under the MP criterion used 500 bootstrap replicates, each with two random taxon addition replicates, and TBR swapping to completion on 10,000 minimal trees found.

Analyses of each dataset using maximum likelihood (ML) were conducted with GARLI v. 0.96b (Zwickl 2006) and with RAxML 7.0.4 (Stamatakis 2006, implemented

on the WWW at the CIPRES Portal 2.0, [www.phylo.org/portal2](http://www.phylo.org/portal2)). GARLI was used to analyze *rbcL* and *matK* independently, while RAxML was used for analysis of the combined data matrix. RAxML allows the estimation of independent substitution models for each data partition rather than estimating a single set of parameters to cover the entire combined matrix. Bootstrap support values for the independent ML analyses of each dataset are determined from 500 GARLI bootstrap replicates. For the combined matrix, 500 rapid bootstrap heuristics replicates from RAxML were executed.

Bayesian analyses were performed on the individual and combined data matrices using MrBayes v.3.1.2 (Huelsenbeck and Ronquist 2001). Models of nucleotide substitution were determined for the *rbcL* and *matK* matrices using the Akaike Information Criteria (AIC) implemented in Modeltest 3.7 (Posada and Crandall 1998). The appropriate form of substitution model (but not the specific estimated model parameter values themselves) was then applied for each data partition in the individual and combined analyses. Prior probability distributions on all parameters were kept at default values, and all model parameters except for tree topology and branch lengths were unlinked between partitions. Two independent analyses were run simultaneously, with the built-in convergence diagnostic (average standard deviation of split frequencies) assuming a relative burn-in of 25%. Analyses were run for a minimum of 4 million MCMC generations (sampling rate = 0.01), at which point they were stopped if the convergence diagnostic had dropped below 0.01. Each Bayesian analysis was repeated to confirm that clade posterior probabilities were stable and repeatable.

**Divergence Time Estimates** — For divergence time estimation, a reduced matrix was utilized that included only the members of Linaceae and Irvingiaceae (*Irvingia malayana*), which was found to be the family most recently diverged from Linaceae in the analyses of Davis et al. (2005). Divergence times were estimated using BEAST 1.4.8 (Drummond and Rambaut 2007) in two separate analyses using the maximum (105 mya) and minimum (94.5 mya) optimal divergence times estimated by Davis et al. (2005) as priors for the age of the root node (the most recent common ancestor of Linaceae and Irvingiaceae). For each analysis, the optimal Linaceae-Irvingiaceae divergence time prior was set to a normal distribution, with the optimal time determined by Davis et al. set as the mean, and a standard deviation of 1.5 (to approximate the uncertainty around the mean reported by Davis et al. [2005]).

A second calibration point for these analyses was based on *Linum* pollen fossils. The oldest known *Linum* pollen grains are from late Eocene Priabonian strata dated approximately 33.9-37.2 mya, from the Ebro Basin of Northeastern Spain (Cavagnetto and Anadón 1996). The pollen grains are tricolpate and appear to be of either the *Linum catharticum* or *Linum austriacum* type described by Punt and Den Breejen (1981), and differ significantly from the pollen described from the Southeast Asian genera of Linoideae (which are pantoporate) and from members of the Hugonioideae (which are tricolporate, often with a reticulate exine patterns; Saad 1962). If accepted as belonging to the genus *Linum*, these fossils seem to provide evidence for the minimum age for the common ancestor of all *Linum* species (including segregate genera) of approximately 33.9 my. This calibration point was implemented in BEAST by setting the *tmrca* prior on

the common ancestor of all *Linum* (including *Cliococca*, *Hesperolinon*, *Radiola*, and *Sclerolinon*) with a lognormal distribution (mean = 0, standard deviation = 1.0, zero offset = 33.9), which allows this divergence to be estimated no later than 33.9 mya, but does allow earlier estimates if supported by the data.

In BEAST, the MCMC chain was run for 30 million generations with the auto-optimization feature activated (initial 10% burn-in discarded from all calculations and estimates), sampling tree topology, model parameters, and divergence times every 1000 generations. The uncorrelated lognormal relaxed clock option was used for sampling branch lengths, with a Yule process model for speciation rate, as recommended in the BEAST manual for estimating divergence times on multispecies phylogenies. At the end of each BEAST run, parameter logs were examined using Tracer v1.4.1 (Rambaut and Drummond) to verify that the MCMC chain had reached stationarity and had sampled divergence time estimates and model parameters an adequate number of times (effective sample sizes > 200, as recommended in the BEAST manual). Because the Hugonioideae were found to be monophyletic in all optimal likelihood-based topologies, the subfamily was constrained to monophyly in the BEAST analysis.

## RESULTS

Sequence and estimated substitution model parameters for each data matrix and the combined matrix are provided in Table 2.6, along with MP and ML-based tree scores obtained in each analysis.



**Missing Data** — Only partial sequences of *rbcL* could be obtained for *Indorouchera griffithiana* and *Philbornea magnifolia* (approximately 400 bp missing from 5' end of the gene). *Hugonia mystax* and *Linum hypericifolium* are similarly missing approximately 400 bp at their 3' ends. *Roucheria monsalveae* and *Linum arboreum* are missing the first 600 bp of *matK*. For *Roucheria calophylla*, the *rbcL* sequence was provided by Dr. Ken Wurdack (Smithsonian National Museum of Natural History); *matK* sequence was not available. The full sequence alignment is provided in Appendix A.

**Results from *rbcL*** — A strict consensus of MP trees is presented in Fig. 2.3, and the ML topology recovered by GARLI is presented in Fig. 2.4 (phylogram in Fig. 2.5). MP, ML, and Bayesian analyses of *rbcL* find both Linaceae and subfamily Linoideae to be monophyletic, with 100% bootstrap (MP and ML) and posterior probability (PP) support. Hugonioideae is shown to be monophyletic in the *rbcL* ML topology, but without significant bootstrap support.

Within the Linoideae, the blue-flowered linums (sections *Linum* and *Dasylinum*) are shown to be a well supported (MP 97%, ML 98%, PP 100%) lineage that is sister to a clade comprising the rest of the subfamily. The Southeast Asian genera are shown as a lineage sister to the yellow-flowered linums (sections *Cathartolinum*, *Linopsis*, and *Syllinum*) and the segregate genera, but this relationship receives low MP bootstrap support (62%) and insignificant (<50%) ML bootstrap and posterior probability.

Within the Hugonioideae, the strict consensus of MP trees and MP bootstrap values show that relationships among the genera and species of this subfamily are not consistently resolved, the only even moderately-supported relationships being the

monophyly of *Durandea* (*Hugonia* sect. *Durandea*, including *Durandea pentagyna*, *D. neocaledonica*, and *H. penicillanthemum* (MP bootstrap 87%), the monophyly of *Hebepetalum* (MP bootstrap 70%), and the sister relationship of *Hugonia obtusifolia* and *H. spicata* (MP bootstrap 90%). The ML bootstrap and Bayesian analysis resulted in similar levels of support for those relationships. In the ML topology, *Hebepetalum* is shown as the earliest-diverging lineage of Hugonioideae. The *Durandea* clade appears sister to *Indorouchera* plus *Philbornea*. *Roucheria* is shown as sister to *Hugonia*, and their lineage as sister to the *Durandea-Indorouchera-Philbornea* clade. None of these intergeneric relationships receives significant ML bootstrap support, however.

Neither MP, ML, or Bayesian analyses of *rbcL* provide support for any family as the sister group to Linaceae. In the MP analysis, relationships among families of Malpighiales are largely unresolved. ML shows Linaceae sister to a clade consisting of representatives of Balanopaceae, Chrysobalanaceae, and Picrodendraceae, similar to the findings of Chase et al. (2002) and Davis and Chase (2004), but without significant bootstrap support. The placement of families of Cronquist's Linales are not well resolved, though they do not appear closely related to Linaceae in the ML topology.

**Results from matK** — The *matK* gene was found to have approximately twice the number of variable sites as *rbcL*. Results from MP and ML analyses of *matK* are summarized in Figs. 2.6 and 2.7, respectively (ML phylogram in Fig. 2.8). The Linaceae and Linoideae are again found to be monophyletic as currently circumscribed, each with 100% bootstrap support under both MP and ML, and 100% posterior probability. Within the Linoideae, *Linum* sections *Linum* and *Dasylinum* (the “blue-flowered” sections) form

a strongly-supported monophyletic group (100% MP and ML bootstraps, 100% PP), and the clade of yellow-flowered *linums* and segregate genera is also well supported (98% MP and ML bootstraps, 100% PP). In the MP analysis of *matK* (Fig. 2.6), *Anisadenia* is shown as sister to the rest of Linoideae (MP bootstrap of 62), while the relationships of *Reinwardtia* and *Tirpitzia* are unresolved with respect to the two lineages of *Linum* and its segregates. The ML topology shows the Southeast Asian genera forming a monophyletic group sister to the rest of the Linoideae, but without significant bootstrap support for their monophyly and with low support (62% ML bootstrap, 75% PP) for their sister relationship to the remainder of Linoideae.

The Hugonioideae is also found to be monophyletic in the *matK* analyses, but with low bootstrap support (73% MP, 55% ML, 59% PP). Within Hugonioideae, *Roucheria* and *Hebepetalum* form a clade (56% MP, 53% ML, 70% PP) that is sister to the Old World genera, which are a well-supported lineage (96% MP, 99% ML, 100% PP). Generic relationships among Old World Hugonioideae are generally well resolved in both analyses of *matK*, with the *Durandea* clade sister to *Indorouchera* and *Philbornea*, and that lineage sister to *Hugonia*.

The *matK* analyses show Linaceae sister to Clusiaceae + Hypericaceae, a result not previously reported, but this relationship is not significantly supported by MP or ML bootstraps analyses. Relationships among the families of Malpighiales are generally not well supported by *matK* alone. Representatives of the families of Linales are shown dispersed throughout the order, not closely-related to Linaceae in the optimal ML topology.

**Results of Combined Analyses** — Parsimony, likelihood, and Bayesian analysis results for the combined data matrix are summarized in Figs. 2.9 and 2.10, with the combined phylogram shown in Fig. 2.11. These analyses also find Linaceae and Linoideae monophyletic (100% MP and ML bootstraps and 100% PP ), and a weakly-supported monophyletic Hugonioideae (57% MP, 52% PP, ML <50%). Within Linoideae, the blue- and yellow-flowered (plus segregates) lineages of *Linum* are found to be well-supported monophyletic groups. The Southeast Asian genera are shown in unresolved relationships in the parsimony analysis, in a polytomy with the yellow-flowered (plus segregates) lineage of *Linum*, with only low bootstrap support (55%) placing them in that clade. In the ML and Bayesian analyses, the Southeast Asian genera form a monophyletic sister group to the rest of Linoideae, with low levels of support (56% ML bootstrap and 52% PP). Parsimony finds Linaceae sister to the Clusiaceae+Hypericaceae, while ML shows Linaceae sister to Phyllanthaceae+Picrodendraceae, both lacking significant support. Members of the Linales are dispersed among the other families of Malpighiales.

**Molecular Dating** — Table 2.6 lists the divergence times obtained for key nodes within Linaceae. The two analyses with different root age priors resulted in very similar age estimates for nodes within Linaceae, with the means generally within 1-2 million years of each other. While the deepest divergence in the family may have originated in the Cretaceous, most of the extant genera and species are estimated to have diverged more recently, with Linoideae beginning to diversify in the Eocene, and Hugonioideae apparently diversifying later, in the Oligocene and Miocene (Fig. 2.12).

## DISCUSSION

**Linaceae is Monophyletic** — In all of our analyses, the Linaceae, in its current accepted circumscription (including representatives all of the genera of both subfamilies), is found to be a strongly supported monophyletic group. This is not a particularly surprising result, although this study is the first to include all of the genera of Linaceae. This study shows that additional sampling from the tropical Linaceae does not result in the phylogenetic aggregation of “Linalean” taxa, with data from two markers. The dispersal of “Linales” families throughout the Malpighiales was presaged by morphological treatments and has been found in other molecular phylogenetic studies that lacked complete sampling of Linaceae. Strong support for family-level relationships in the order is generally lacking, but our analysis of the combined data matrix find some support for sister-group relationships of some “Linales” families with families other than Linaceae: Erythroxylaceae (represented by *Erythroxylum novocaledonicum*) to Rhizophoraceae (*Bruguiera gymnorhiza*) and Ixonanthaceae (*Ixonanthes reticulata*) to Ochnaceae (*Gomphia serrata*). The phylogenetic positions of Ctenolophonaceae and Humiriaceae are not well resolved. Recent studies of Malpighiales phylogeny with data from 17 chloroplast, nuclear, and mitochondrial markers also fail to resolve relationships among many families of the order, and do not identify the sister family of Linaceae, but do show the families of “Linales” to be dispersed among several different lineages (Wurdack and Davis 2009).

**Subfamilial Relationships** — While Linoideae is found to be a well-supported monophyletic grouping in all of our analyses, the Hugonioideae receives only tepid support in the analyses of *matK* and the combined matrix. This seems to stem from the uncertain placement of the New World Hugonioideae, *Roucheria* and *Hebepetalum*. Our analyses tend to show the South American taxa as a distinct lineage, with the split between Old and New World Hugonioideae occurring at the base of the subfamily. Phylograms of the optimal *rbcL* and *matK* topologies (Figs. 2.5, 2.8) show that the branches at the base of both subfamilies are very short. This may be indicative of relatively rapid divergence within each subfamily (Fig. 2.12), making relationships among their earliest-diverging lineages difficult to resolve. While the molecular data tend to ally *Roucheria* and *Hebepetalum* with the Old World Hugonioideae, they do bear some morphological distinctions from the Old World Genera: their entirely arborescent habit and complete lack of any kind of hooked tendril or other organ that facilitates a climbing habit. Collection of additional data from additional chloroplast and other markers, as well as additional taxa, particularly from *Roucheria* and *Hugonia*, will be necessary to clarify this situation.

Since both subfamilies are shown to be monophyletic in the optimal topologies (except for *rbcL* under MP criteria, Fig. 2.3), our results do not prohibit the recognition of Hugoniaceae as a distinct family, sister to the Linaceae s.s. This represents a departure from the results of McDill et al. (2009), who found Hugonioideae to be paraphyletic with more limited sampling. Their sampling of Hugonioideae had the primary goal of rooting the Linoideae phylogeny, not examining the monophyly of Hugonioideae. While the

paraphyly of Hugonioideae in that earlier study may have resulted from the limited sampling from Hugonioid taxa and outgroups, the lack of statistical support for Hugonioid monophyly in the current analyses leaves the matter open for further investigation.

**Generic Relationships in Hugonioideae** — With the exception of the New World lineage as described above, relationships within the Hugonioideae are resolved fairly consistently. The *Durandea* clade is a consistently found monophyletic group, sister to the small segregate genera *Philbornea* and *Indorouchera*, which has implications for the systematics of *Hugonia*. *Philbornea* and *Indorouchera* have usually been maintained as distinct from *Hugonia* due to their entirely glabrous surface (*Hugonia* species are generally pubescent, at least on the calyx (Van Hooren and Nootboom 1984), different inflorescence structure (axillary racemes or fascicles versus panicles in *Hugonia*), and hook morphology (“hooked branchlets” in *Philbornea* and *Indorouchera*, respectively, versus “hooked tendrils” in *Hugonia* (Jardim 1999)). *Philbornea* and *Indorouchera* have also been maintained as distinct from each other, due to differences in inflorescence structure and the presence of resin on the buds and stipules of *Indorouchera* (van Hooren and Nootboom 1984). *Durandea*, though, has received inconsistent treatment, either as a section of *Hugonia* or a distinct genus. In their 1984 treatment for Flora Malesiana, Van Hooren and Nootboom, treating *Durandea* as a section of *Hugonia*, note that it is distinguished from sect. *Hugonia* (which has a notable indumentum) by being nearly glabrous on all parts (like *Indorouchera* and *Philbornea*), and having a fruit that splits into five pyrenes (the fruit is a coherent drupe-like berry in

most species of *Hugonia* sect. *Hugonia* and in *Indorouchera* and *Philbornea*). *Durandea* also differs from *Hugonia* in lacking glands between the petals, as noted by Jardim (1999), who considered *Durandea* as a genus. Saad (1962) noted similarities in the pollen morphologies of *Durandea* and *Philbornea*.

Our analyses show that the *Durandea* clade, *Indorouchera*, and *Philbornea* constitute a lineage that is sister to the rest of *Hugonia*. *Hugonia penicillanthemum* appears to be part of the *Durandea* clade; although this species apparently has never been treated as a species of *Durandea*, its morphology indicates it is a member of *Hugonia* sect. *Durandea*. While it would be premature to conclude definitively that *Durandea* is a monophyletic genus distinct from *Hugonia* (since additional sampling from either taxon could alter our understanding of their relationship), it seems apparent that *Durandea* probably should not be treated as a section of *Hugonia* unless *Indorouchera* and *Philbornea* are placed in *Hugonia* as well. Additional revisionary studies of the Hugonioideae, and increased phylogenetic sampling, will be necessary to address these questions of generic taxonomy.

**Generic Relationships in Linoideae** — With respect to relationships within Linoideae, the results of McDill et al. (2009) are largely compatible with those found here, with minor differences perhaps due to different taxon sampling and data. Their results also do not strongly resolve the position of the Southeast Asian genera. Their taxonomic sample included only a single species from each genus, whereas here we include three (of three) *Anisadenia* and two (of three) *Tirpitzia*, finding both genera to be monophyletic and well-supported.



**Divergence Times and Biogeography** — The divergence times estimated in this study (Table 3.6) also depart significantly in some cases from those estimated in McDill et al. (their Table 4) for nodes the two studies have in common. The estimates of McDill et al. are generally older, falling near the upper (older) bound of the 95% HDP confidence intervals estimated in BEAST. This may be a result of altered taxon sampling and additional data in the present analysis, the use here of an additional fossil calibration point, and the difference in estimation method (McDill et al. employed penalized likelihood conditioned on the optimal ML *rbcL* topology obtained in their Linoideae-centric analyses, whereas we here employ a Bayesian methodology sampling different potential topologies and branch lengths).

Davis et al. (2005) concluded that the order Malpighiales underwent a rapid radiation in the late Aptian, from 94 to 114 million years ago, as elements of moist, closed-canopy, megathermal rain forests. The earliest common ancestor of Linaceae may indeed have originated within that time frame (node A), but our results show that the extant tropical components of Linaceae, the Hugonioideae, diversified at a much later time, with their earliest divergence occurring between 30 and 11 mya, with the average of divergence time estimates of that clade in the early Miocene (Fig. 2.12, node I). The Miocene is thought to have exhibited a relatively warm global climate optimal for rainforest expansion (Zachos et al. 2001, Morley 2000), which could have influenced the diversification of extant tropical Linaceae. The Linoideae appears to have initially diversified in the mid-to-late Eocene (Fig. 2.12, node C), a period of cooling global climate that promoted the establishment of communities dominated by grasses and other

herbaceous taxa in the Northern Hemisphere (Tiffney and Manchester 2001), habitats where species of *Linum* and its segregates are commonly found today.

The Southeast Asian tropical/subtropical genera of Linoideae are shown to be the basally-diverging members of that subfamily, with moderate ML bootstrap and Bayesian support (Fig. 2.10). This may indicate that the more temperate members of Linaceae (*Linum* and its segregates) are derived from tropical or subtropical ancestry, but not from within the extant, tropical subfamily Hugonioideae. In the absence of a definitive sister-group for Linaceae, the characteristics – ecological or morphological – of the common ancestor of Linoideae and Hugonioideae remain a mystery. Interpretation of the biogeographic history of the Hugonioideae is further hindered by the lack of strongly supported basal resolution in the subfamily. With the current taxon sampling and phylogenetic resolution, and given the lack of a consistently identified outgroup for the Linaceae, it is not possible to determine whether the Hugonioideae, or Linaceae in general, had its origins in the Old World or New World tropics.

The presence of Hugonioideae in the tropics of Africa and South America, with the basal phylogenetic split in the subfamily occurring perhaps between Old World and New World lineages, brings to mind the biogeographic pattern of Gondwanan vicariance (Raven and Axelrod 1974). However, considering published dates for the break-up of Gondwana (McLoughlin 2001), a Gondwanan biogeographic scenario would require the most recent common ancestor of these taxa to be present prior to the separation of South America and Africa, which was complete by approximately 100 million years ago. The Hugonioideae seem to have diverged much more recently, approximately 40 million

years ago at the earliest, too late to have been affected by the breakup of Gondwana. *Hebepetalum* and *Roucheria* species are estimated to have diverged from from Old World ancestry within the last 10 mya (Fig. 2.12), which would also seem to preclude a “Boreotropical” scenario involving the land bridge thought to have spanned the North Atlantic during the middle and late Eocene (Tiffney 1985). This land bridge is thought to have been concurrent with a global thermal maximum that could have allowed migration of tropical taxa at relatively high latitudes (Tiffney and Manchester, 2001). Such scenarios have been found to be compatible with the phylogenies, fossil distributions, and divergence times of several angiosperm lineages that currently exhibit pantropical distributions similar to Hugonioideae, such as the Burseraceae (Weeks et al. 2005) and the Malpighiaceae (Davis et al. 2002). Long-distance dispersals have also been implicated as important factors in the establishment of Northern and Southern Hemisphere disjunct distributions in plants (Givnish and Renner, 2004; Pennington et al. 2004; Sanmartin and Ronquist, 2004), and may have played a role in biogeographic history of Linaceae, but adequate testing of these hypotheses in Hugonioideae awaits more robust phylogenetic resolution and more extensive taxon sampling.

## CONCLUSION

Our results indicate that the two subfamilies of Linaceae are monophyletic and sister to each other, although the monophyly of Hugonioideae is not well supported. Our results do not indicate that any members of Cronquist’s Linales, or the historical, expanded concepts of Linaceae, are closely related to Linaceae, but support for

relationships among families in Malpighiales is paltry. We can conclude that *Durandea* might best be recognized as a genus distinct from *Hugonia* rather than a section of it, unless the species currently placed in *Indorouchera* and *Philbornea* are transferred to *Hugonia* as well. A global taxonomic revision of *Durandea* and *Hugonia* is recommended in order to clarify generic circumscriptions and membership. Based on estimates of divergence times, the biogeographic pattern exhibited by the Hugonioideae does not seem to be the result of Gondwanan vicariance or Boreotropical migration, which implicates long-distance dispersal as a remaining possible mechanism for the establishment of their pantropical distribution.

Table 2.1. The genera of Linaceae, with the number of species known in each, the number sampled for this investigation of subfamilial relationships, and their general geographic distributions.

	# of species (# sampled)	Distribution
<b>Subf. Linoideae</b>		
<i>Anisadenia</i>	3(3)	Southeast Asia
<i>Cliococca</i>	1(1)	Temperate South America
<i>Hesperolinon</i>	13(4)	W. North America
<i>Linum</i>	187(17)	Temperate Cosmopolitan
<i>Radiola</i>	1(1)	Europe
<i>Reinwardtia</i>	2(1)	Southeast Asia
<i>Sclerolinon</i>	1(1)	W. North America
<i>Tirpitzia</i>	3(2)	Southeast Asia
<b>Subf. Hugonioideae</b>		
<i>Durandea</i>	15(3)	Malaysia
<i>Hebepetalum</i>	3 (2)	Tropical S. America
<i>Hugonia</i>	30 (7)	Tropical Africa, SE Asia
<i>Indorouchera</i>	2(1)	Indonesia
<i>Philbornea</i>	1(1)	Indonesia
<i>Roucheria</i>	7 (3)	Tropical S. America

Table 2.2. Historical treatments of Linaceae, and the current, generally accepted treatment (Dressler et al. in press).

<b>Planchon 1848</b>	<b>Bentham and Hooker 1862</b>	<b>Baillon 1878</b>	<b>Winkler 1931</b>	<b>Dressler et al.</b>
		<b>Subf. Lineae</b>	<b>Subf. Linoideae</b>	<b>Subf. Linoideae</b>
<b>Sect. Eulineae</b>	<b>Tribe Eulineae</b>		<b>Tribe Eulineae</b>	
<i>Linum</i>	<i>Linum</i>	<i>Linum</i>	<i>Linum</i>	<i>Linum</i>
<i>Radiola</i>	<i>Radiola</i>	<i>Radiola</i>	<i>Radiola</i>	<i>Radiola</i>
<i>Reinwardtia</i>	<i>Reinwardtia</i>		<i>Reinwardtia</i>	<i>Reinwardtia</i>
			<i>Tirpitzia</i>	<i>Tirpitzia</i>
			<i>Hesperolinon</i>	<i>Hesperolinon</i>
				<i>Sclerolinon</i>
<b>Sect. Anisadeneae</b>			<b>Tribe Anisadenieae</b>	<i>Cliococca</i>
<i>Anisadenia</i>	<i>Anisadenia</i>	<i>Anisadenia</i>	<i>Anisadenia</i>	<i>Anisadenia</i>
<b>Sect. Hugonieae</b>	<b>Tribe Hugonieae</b>	<b>Subf. Hugonieae</b>	<b>Tribe Hugonieae</b>	<b>Subf. Hugonioideae</b>
<i>Hugonia</i>	<i>Hugonia</i>	<i>Hugonia</i>	<i>Hugonia</i>	<i>Hugonia</i>
<i>Roucheria</i>	<i>Roucheria</i>		<i>Roucheria</i>	<i>Roucheria</i>
<i>Durandea</i>			<i>Durandea</i>	<i>Durandea</i>
	<b>Tribe Erythroxyleae</b>	<b>Subf. Erythroxyleae</b>	<i>Philbornea</i>	<i>Philbornea</i>
	<i>Erythroxylum</i>	<i>Erythroxylum</i>	<i>Indorouchera</i>	<i>Indorouchera</i>
	<i>Hebepetalum</i>		<i>Hebepetalum</i>	<i>Hebepetalum</i>
	<i>Aneulophus</i>	<i>Aneulophus</i>		
	<b>Tribe Ixonanthese</b>		<b>Subf. Ixonanthoideae</b>	
	<i>Ochthocosmus</i>	<i>Ochthocosmus</i>	<i>Ochthocosmus</i>	
	<i>Ixonanthes</i>	<i>Ixonanthes</i>	<i>Ixonanthes</i>	
	<i>Durandea</i>		<b>Subfamily Ctenolophonoideae</b>	
	<i>Phyllocosmus</i>		<i>Ctenolophon</i>	
	<i>Sarcotheca</i>			
		<b>Subfamily Houmirieae</b>	<b>Subf. Humirioideae</b>	
		<i>Humiria</i>	<i>Humiria</i>	
			<i>Sacoglottis</i>	
			<i>Vantanea</i>	
			<b>Tribe Nectaropetaleae</b>	
			<i>Peglerya</i>	
			<i>Nectaropetalum</i>	

Table 2.3. Voucher specimens for taxa sampled for *matK* and *rbcL*, and GenBank accession numbers for previously published sequences included in analyses. For new sequences obtained for this study, GenBank Accession numbers are designated as Tba (“to be announced”), and will be obtained upon journal publication. Sequence alignments are provided in Appendix A.

Taxon	Distribution	Voucher Collection	Herb.	Locality	Genbank matK	Accession # rbcL
<b>Linoideae</b>						
<i>Anisadenia khasayana</i> Griff.	Nepal	F. Miyamoto et al. 9440158	GH	Nepal	Tba	tba
<i>Anisadenia pubescens</i> Griff.	China	Bartholomew et al. 1984 Sino-American Bot. Exped. No. 1011	GH	China	Tba	FJ169557
<i>Anisadenia saxatilis</i> Wall. Ex Meissn.	Nepal	M. Mikage et al. 9470573	GH	Nepal	Tba	tba
<i>Cliococca selaginoides</i> (Lam.) C.M.Rogers & Mildner	S. Am.	A. Krapovickas & C.L. Cristóbal 34169	MO	Brazil,	TBA	FJ169558
<i>Hesperolinon clelandii</i> Small	N. Am	Ertter 8537	TEX	CA	Tba	tba
<i>Hesperolinon disjunctum</i> Sharsm.	N. Am.	McDill 87	TEX	CA	Tba	tba
<i>Hesperolinon drymarioides</i> Small	N. Am	Dibble and Griggs 14	MO	CA	tba	FJ169560
<i>Hesperolinon micranthum</i> Small	N. Am.	H. van der Werff and D. Clark 4574	MO	CA	TBA	FJ169561
<i>Linum arboreum</i> L.	Europe	Dechamps 9103	MO	Greece	Tba	tba
<i>Linum berlandieri</i> Hook.	N. Am.	B.L. Turner 21-427	TEX	TX, Kerr Co.	TBA	FJ169567
<i>Linum bienne</i> Mill.	Europe	McDill 2004-XII-5.1	TEX	Chile (introduced)	Tba	tba
<i>Linum catharticum</i> L.	Europe	A.K. Skvortsov et al. 4263078	MO	Estonia	Tba	tba
<i>Linum comptonii</i> C.M. Rogers	S. Africa	Elsie Esterhuysen 35882	MO	South Africa	TBA	FJ169572
<i>Linum flavum</i> L.	Eurasia	C.M. Rogers 13374	MO	Turkey	TBA	TBA
<i>Linum hirsutum</i> L.	Eurasia	P. Frost-Olsen 4542	MO	Bulgaria	Tba	tba
<i>Linum hypericifolium</i> Salisb.	Eurasia	M. Merello et al. 2116	MO	Georgia	Tba	tba
<i>Linum lewisii</i> Pursh	N. Amer.	Turner 21-263	TEX	TX, Reeves Co.	n.a.	tba
<i>Linum lewisii</i> Pursh		n.a.	n.a.	n.a.	AB233791	n.a.
<i>Linum monogynum</i> G. Forst.	Australia	Meudt 245	TEX	Tasmania	Tba	tba
<i>Linum narbonense</i> L.	Europe	C. Ardo et al. 2303	TEX	Spain	Tba	tba
<i>Linum nervosum</i> Waldst. & Kit.	Eurasia	MBG 4250021	MO	Russia	Tba	tba
<i>Linum oligophyllum</i> Willd.	S. Am.	D.N. Smith et al. 9426	MO	Peru	TBA	TBA
<i>Linum perenne</i> L.	Eurasia	MJG 040929	MJG	(cultivated)	na	FJ169582
<i>Linum perenne</i> L.		n.a.	n.a.	n.a.	AB038182	na
<i>Linum pratense</i> Small	N. Am	McDill 2005-V-13.1	TEX	TX, Montague Co.	tba	tba
<i>Linum rupestre</i> (A.Gray) Engelm. ex A.Gray	N. Am.	B.L. Turner 21-589	TEX	TX, Sutton Co.	TBA	FJ169586
<i>Linum suffruticosum</i> L.	Europe	C. Navarro et al. CN 2339	TEX	Spain	Tba	FJ169591
<i>Linum usitatissimum</i> L.	Eurasia	MJG 040924	MJG	(cultivated)	Tba	FJ169596
<i>Linum volkensii</i> L.	E. Africa	R.E. Gereau & C.J. Kayombo 4662	MO	Tanzania	Tba	FJ169597
<i>Radiola linoides</i> Roth.	Eurasia, N.	Knut Egeröd 6935	MO	Sweden	Tba	tba

<i>Reinwardtia indica</i> Dum.	Africa Nepal, India	McDill s.n. 2002	TEX	(cultivated)	tba	FJ169599
<i>Sclerolinon digynum</i> (A. Gray) C.M. Rogers	N. Am.	M.S. Taylor 3935	MO	CA, Plumas Co.	TBA	FJ169601
<i>Tirpitzia ovoidea</i> Chun & F.C.Ho	China	Wei Fanan 1806	MO	China	Tba	tba
<i>Tirpitzia sinensis</i> (Hesml.) Hall.	China	Libo Team 81-6-0105	MO	China	tba	tba
<b>Hugonioideae</b>						
<b><i>Durandea</i> (=Hugonia subg. <i>Durandea</i>)</b>						
<i>Durandea pentagyna</i> K. Schum	SW Pacific	W. Takeuchi 4523	MO	Papua New Guinea	tba	FJ169559
<i>Durandea neocaledonica</i> (Vieill.) Baum.-Bod.	New Caledonia	Gordon McPherson 5443	MO	New Caledonia	Tba	tba
<i>Hugonia penicillanthemum</i> Baill. ex Pancher & Sebert	W Pacific	A. Gentry & G. McPherson 34683	MO	New Caledonia	Tba	tba
<b><i>Hebepetalum</i></b>						
<i>Hebepetalum neblinae</i> A. Jardim & P.E. Berry	S. Amer.	P. E. Berry, C. Gómez, D. Acosta 6319	MO	Venezuela	Tba	tba
<i>Hebepetalum humirifolium</i> Benth.	S. Amer.	J.J. Wurdack 2467	F	Peru	Tba	tba
<b><i>Hugonia</i> (=Hugonia subg. <i>Hugonia</i>)</b>						
<i>Hugonia</i> aff. <i>planchonii</i> Hook. F.	Africa	Gordon McPherson 16804	MO	Gabon	Tba	tba
<i>Hugonia busseana</i> Engl.	Africa	A. Chikuni, I.H. Patel, W. Nachamba 144	MO	Malawi	Tba	tba
<i>Hugonia castaneifolia</i> Engl.	Africa	R.E. Gereau et. al 6206	MO	Tanzania	Tba	tba
<i>Hugonia gabunensis</i> Engl.	Africa	Lee White 0908	MO	Gabon	Tba	FJ169563
<i>Hugonia longipes</i> H. Perr.	Africa	G.E. Schatz & J.S. Miller 2476	MO	Madagascar	Tba	tba
<i>Hugonia mystax</i> L.	S.E. Asia	P.L. Comanor 414	MO	Sri Lanka	Tba	tba
<i>Hugonia obtusifolia</i> C.H. Wright	Africa	Lee White 0978	MO	Gabon	Tba	tba
<i>Hugonia orientalis</i> Engl.	Africa	P. van Wyk #BSA 2606	MO	Zimbabwe	Tba	tba
<i>Hugonia spicata</i> Oliv.	Africa	Carvalho 3850	MO	Equatorial Guinea	Tba	tba
<b><i>Indorouчера</i></b>						
<i>Indorouчера griffithana</i> (Planch.) Hallier f.	SE Asia	T.G. Laman et al. TL 1155	GH	Indonesia	Tba	tba
<b><i>Philbornea</i></b>						
<i>Philbornea magnifolia</i> Hallier f.	W Pacific	Niga Nangkat NN 161	GH	Brunei	Tba	tba
<b><i>Rouчерia</i></b>						
<i>Rouчерia calophylla</i> Planch.	S. Amer.	Alencar 593	NY	Brazil	n.a.	FJ169600
<i>Rouчерia monsalvae</i> A.H. Gentry	S. Amer.	Miryam Monsalve B. 1996	MO	Colombia	tba	tba
<i>Rouчерia schomburgkii</i> Planch.	S. Amer.	Manuel Rimachi Y. 8416	MO	Peru	tba	FJ169603
<b>Other Malpighiales</b>						
<i>Campostylus mannii</i> (Oliver) Gilg.	Achariaceae				AB233736	AB233840
<i>Kiggelaria africana</i> L.	Achariaceae				AB233739	AB233843
<i>Balanops pancheri</i> Baill.	Balanopaceae				AB233740	AB233844
<i>Euphronia guianensis</i> (R.H. Schomb.) Hallier f.	Chrysobalanaceae				AB233741	AB233845
<i>Trigonía boliviana</i> Warm.	Chrysobalanaceae				AB233744	AB233848
<i>Clusia rosea</i> Jacq.	Clusiaceae				AB233745	AB233849
<i>Garcinia subelliptica</i> Merr.	Clusiaceae				AB233746	AB233850
<i>Ctenolophon englerianus</i>	Ctenolophonaceae				EF135524	AJ402940



Mildbr.			
<i>Elatine triandra</i> Schkuhr	Elatinaceae	AB233749	AB233853
<i>Erythroxylum novocaledonicum</i> O.E. Schulz	Erythroxylaceae	AB233826	AB233930
<i>Acalypha insulana</i> Müll. Arg.	Euphorbiaceae	AB233750	AB233854
<i>Euphorbia humifusa</i> Willd.	Euphorbiaceae	AB233780	AB233884
<i>Manihot esculenta</i> Crantz.	Euphorbiaceae	AB233776	AB233880
<i>Pimelodendron griffithianum</i> (Müll.Arg.) Benth.	Euphorbiaceae	AB233783	AB233887
<i>Ricinus communis</i> L.	Euphorbiaceae	AB233767	AB233871
<i>Jatropha integerrima</i> Jacq.	Euphorbiaceae	AB233775	AB233879
<i>Humiria balsamifera</i> Aubl.	Humiriaceae	AB233785	AB233889
<i>Sacoglottis</i> sp.	Humiriaceae	AB233786	AB233890
<i>Cratoxylum cochinchinense</i> Blume	Hypericaceae	AB233787	AB233891
<i>Irvingia malayana</i> Oliver	Irvingiaceae	AB233789	AB233893
<i>Ixonanthes reticulata</i> Jack	Ixonanthaceae	AB233789	AB233893
<i>Lacistema aggregatum</i> (P.J. Bergius) Rusby	Lacistemaceae	AB233790	AB233894
<i>Byrsonima crassifolia</i> H.B.&K.	Malpighiaceae	AB233794	AB233898
<i>Malpighia glabra</i> L.	Malpighiaceae	AB233796	AB233900
<i>Gomphia serrata</i> (Gaertn.) Kanis	Ochnaceae	AB233803	AB233907
<i>Passiflora quadrangularis</i> L.	Passifloraceae	AB233808	AB233912
<i>Bischofia javanica</i> Blume	Phyllanthaceae	AB233813	AB233917
<i>Flueggea virosa</i> (Roxb. Ex Willd.) Voigt	Phyllanthaceae	AB233815	AB233919
<i>Phyllanthus flexuosus</i> (Siebold & Zucc.) Müll. Arg.	Phyllanthaceae	AB233817	AB233921
<i>Scagea oligostemon</i> (Guillaumin) McPherson	Picrodendraceae	AB233821	AB233925
<i>Drypetes littoralis</i> (C.B. Rob.) Merr.	Putranjivaceae	AB233823	AB233927
<i>Bruguiera gymnorhiza</i> (L.) Lam.	Rhizophoraceae	AB233823	AB233927
<i>Flacourtia indica</i> Merr.	Salicaceae	AB233829	AB233933
<i>Lunania parviflora</i> Spruce ex. Benth.	Salicaceae	AB233832	AB233936
<i>Ionidium commune</i> A.St.-Hil.	Violaceae	AB233836	AB233940
<b>Outgroups</b>			
<i>Euonymus sieboldianus</i> Blume	Celastraceae	AB233838	AB233942
<i>Datisca cannabina</i> L.	Datisceae	AB016467	AB016453
<i>Albizia tomentosa</i> Standl.	Fabaceae	AF5234093	n.a.
<i>Albizia julibrissin</i> Durazz.	Fabaceae	n.a.	Z70147
<i>Quercus rubra</i> L.	Fagaceae	AY312058	AB125026
<i>Oxalis corniculata</i> L.	Oxalidaceae	AB233839	AB233943
<i>Kerria japonica</i> (L.) DC.	Rosaceae	AB073686	AF132893

Table 2.4. Primer sequences.

Primer name	Sequence (5'-3')	Reference
rbcL 1F	ATGTCACCACAAACAGAAAC	Olmstead et al. 1992
rbcL 507F	TAYTGGGCTGTACTATTAAAC	McDill et al. 2009
rbcL 507R	GTTTAATAGTACAGCCCA	(rev/comp of 507F)
rbcL 4R Hugonioid	GTCTATCAATAACTGCATGCATTGC	This study.
rbcL 4F Hugonioid	GCAATGCATGCAGTTATTGATAGAC	This study.
rbcL 1460 R	CTTTTAGTAAAAGATTGGGCCGAG	Chase et al. 1993
rbcL Rb Hugonioid	CTCCTTCCATAYTTCACAAGCAGC	This study.
matK 1F	ACTGTATCGCACTATGTATCA	Sang et al. 1997
matK 1320R	GATCCGCTATAATAATGAGA	Sang et al. 1997
HugoB	CTTTCGCTCAAGAAGGAACCGA	This study
BlueA	TAGCTTYTTCTTGAGCGAAAT	This study
BlueB	ATTTCGCTCAAGAARRAAGCT	This study
648F Linum	CAAGCGGGATCTTTTATCCA	This study
648R Linum	TGGATAAAAGATCCCGCTTG	This study
425F Linum	CTTTCGCTCAAGAAGAACTG	This study
425R Linum	TCAGTTTCTTCTTGAGCGAAAG	This study

Table 2.5. Sequence information.

	rbcL	matK	Combined
Number of Taxa (Linaceae only)	92 (51)	91 (50)	92 (51)
Aligned Length	1293	1504	2797
Included characters	1293	1305	2586
Constant characters (Linaceae only)	845 (1112)	444 (871)	1289 (1983)
Variable characters (Linaceae only)	448 (181)	861 (434)	1309 (615)
Informative characters (Linaceae only)	305 (113)	659 (307)	964 (420)
# of MPT recovered	100158	1898	314
MPT Length (inf. only)	1725 (1566)	3691 (3440)	5491
CI (inf. only)	0.358 (0.293)	0.411 (0.368)	0.389 (0.340)
RI	0.671	0.709	0.690
RC (inf only)	0.241 (0.197)	0.291 (0.261)	0.268 (0.234)
Substitution Model Selected (AIC)	GTR+I+G	TVM+I+G	GTR+I+G
freqA	0.2665	0.3118	0.2936
freq C	0.1784	0.1564	0.1697
Freq G	0.2504	0.1447	0.1912
Freq T	0.3046	0.3871	0.3455
[A-C]	1.3285	1.670	1.2579
[A-G]	3.0656	2.1976	2.3599
[A-T]	0.6041	0.2090	0.3270
[C-G]	0.9496	1.1763	0.9925
[C-T]	4.8131	2.1976	2.9488
[G-T]	1.000	1.0000	1.0000
Gamma Shape	0.7312	1.3508	1.1402
Proportion of Invariable Sites	0.5422	0.0980	0.3652
-Ln L (modeltest)	10883.4316	19776.4453	31400.4297
-Ln L (GARLI, RAxML for combined)	10785.7401	19706.8311	31221.1671

Table 2.6. Estimated ages for nodes indicated in Fig. 2.13, in millions of years before present. Mean divergence times and confidence interval (95% height density proportion) estimated in BEAST, with the age of the root set at the maximum and minimum optimal values estimated by Davis et al. (2005) based on combined atpB, rbcL, 18S, and nad1B-C sequences and several fossil calibrations.

MRCA of:	Root at 105 mya			Root at 94.5 mya		
	95% HPD upper	mean	95% HPD lower	95% HPD upper	mean	95% HPD lower
A) Linaceae and Irvingiaceae	106.83	104.88	102.92	96.36	94.39	92.48
B) Linaceae	80.90	58.77	41.00	76.53	56.05	40.42
C) Linoideae	47.90	37.94	33.93	43.96	36.97	33.93
D) <i>Linum</i> (including segregate genera)	46.79	37.31	33.92	42.85	36.46	33.92
E) <i>Anisadenia</i> , <i>Reinwardtia</i> , <i>Tirpitzia</i>	44.55	30.46	15.50	41.26	29.35	15.41
F) <i>Radiola</i> and yellow-flowered <i>Linum</i>	38.43	29.01	20.54	35.78	28.20	20.18
G) Blue-flowered <i>Linum</i>	35.85	26.71	17.79	33.66	25.90	17.98
H) Yellow-flowered <i>Linum</i>	32.46	24.50	16.58	30.96	23.77	16.35
I) Hugonioideae	37.59	22.83	11.53	35.35	21.84	10.96
J) <i>Linum perenne</i> and <i>L. nervosum</i>	25.53	16.80	8.05	24.63	16.20	8.39
K) North American Blue-flowered <i>Linum</i>	3.70	1.42	0.02	3.62	1.38	0.01

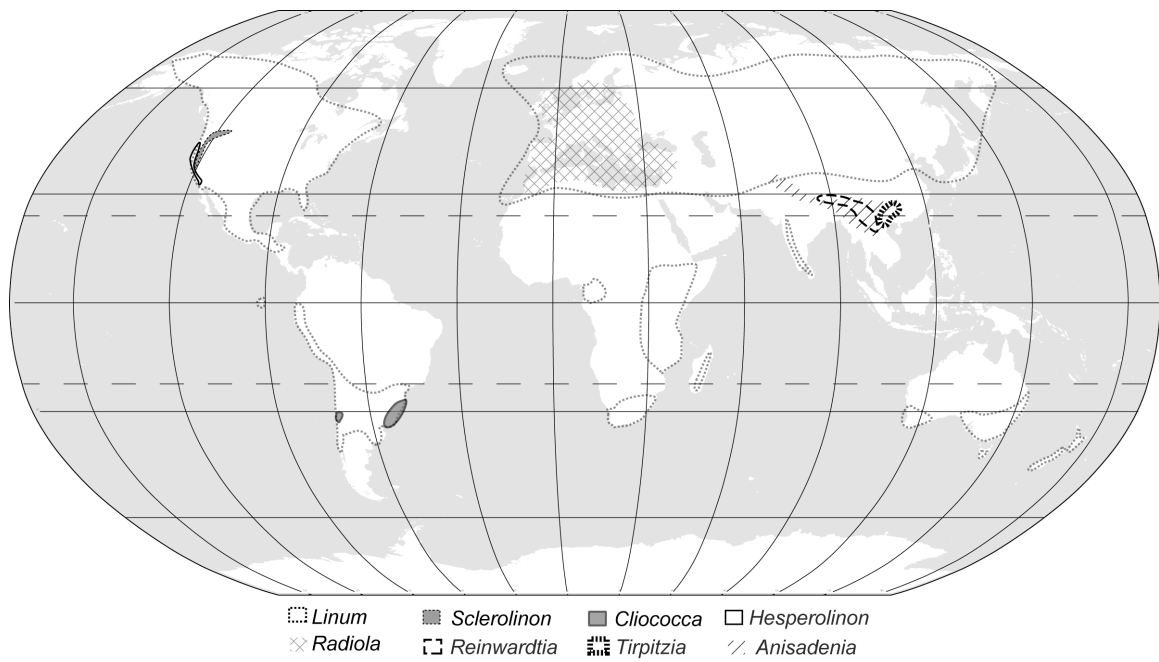


Fig. 2.1: Distributions of the eight genera of Linaceae subf. Linoideae.

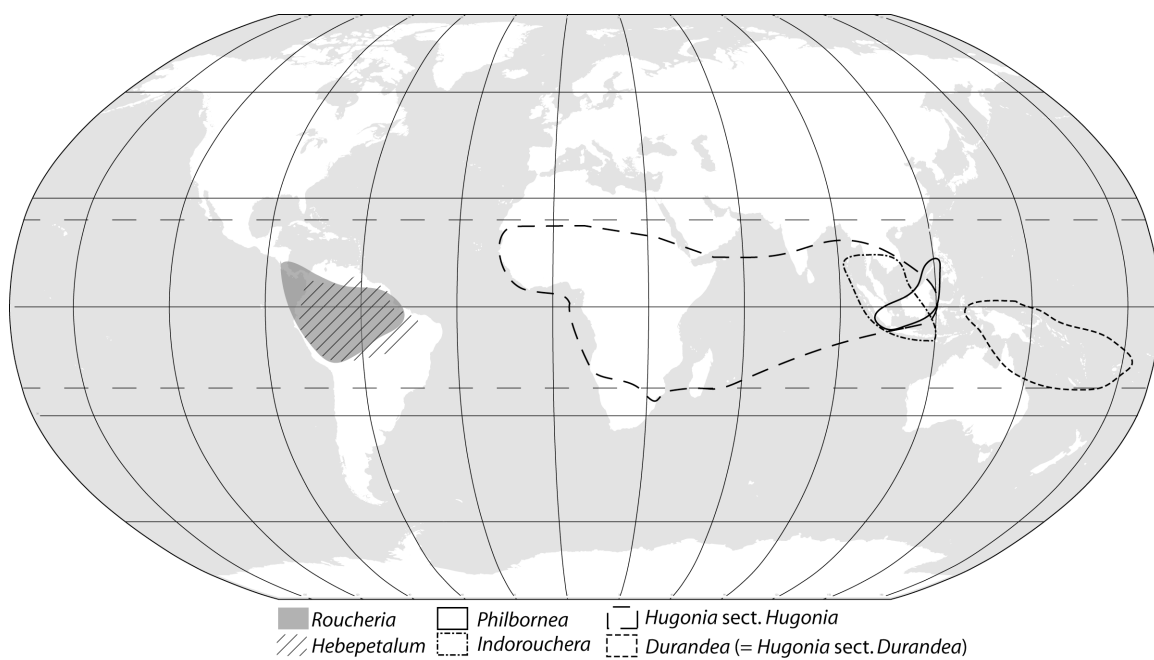


Fig. 2.2. Distributions of the six genera of Linaceae subfamily Hugonioideae, based on Jardim (1999) and Van Hooren and Nootboom (1984).

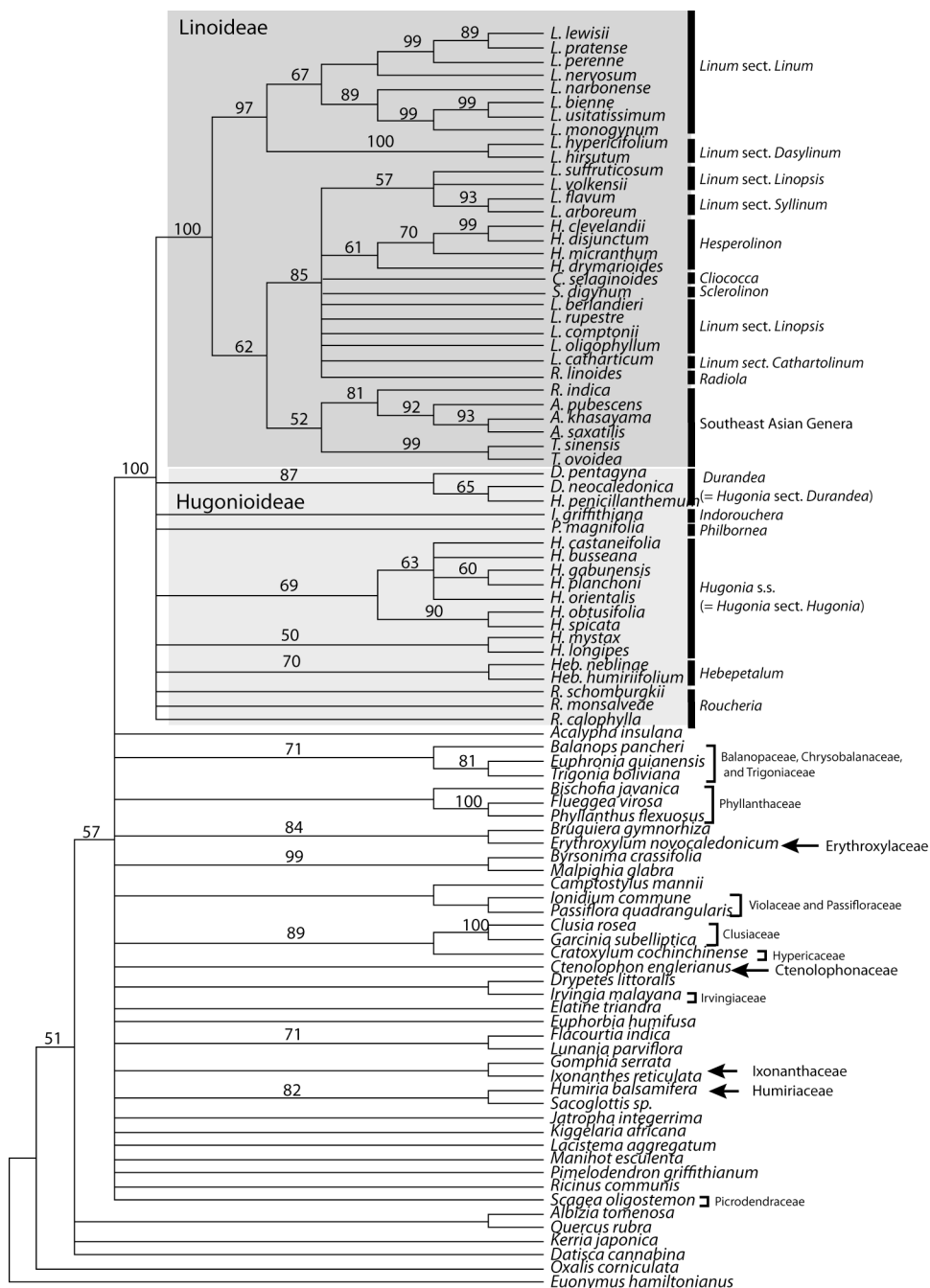


Fig. 2.3. Strict consensus of 100, 158 MP trees (L=1725) based on rbcL data only, with MP bootstrap values above branches (<50% not shown). Linaceae subfamilies as indicated by shaded boxes, genera by black bars. Arrows indicate representatives of the families of Cronquist's Linales, brackets taxa that have appeared in sister-lineages to the Linaceae in previous studies.

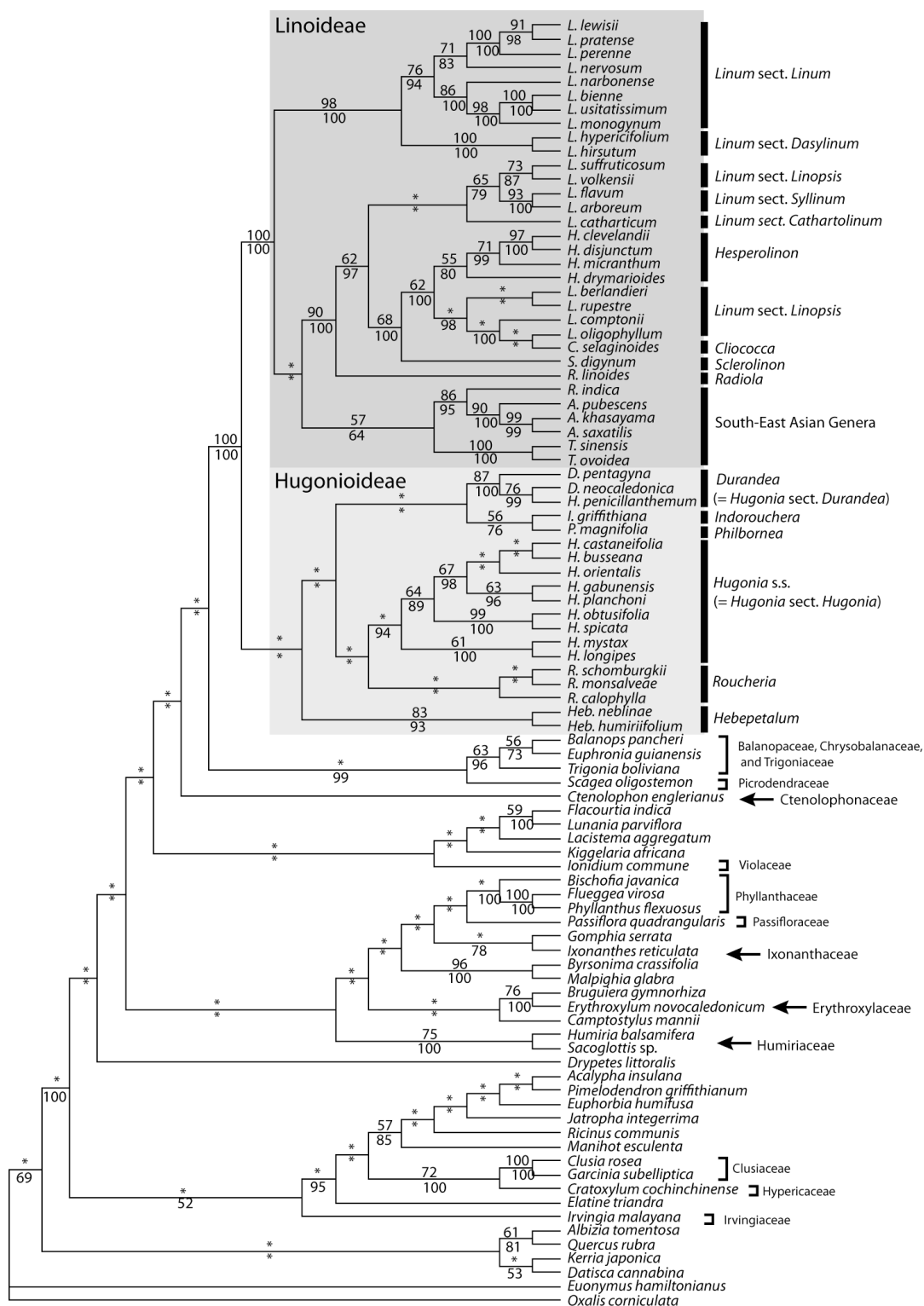


Fig. 2.4: ML topology for rbcL recovered by GARLI, with ML bootstrap percentages above branches, Bayesian posterior probabilities below (values <50% indicated by \*).



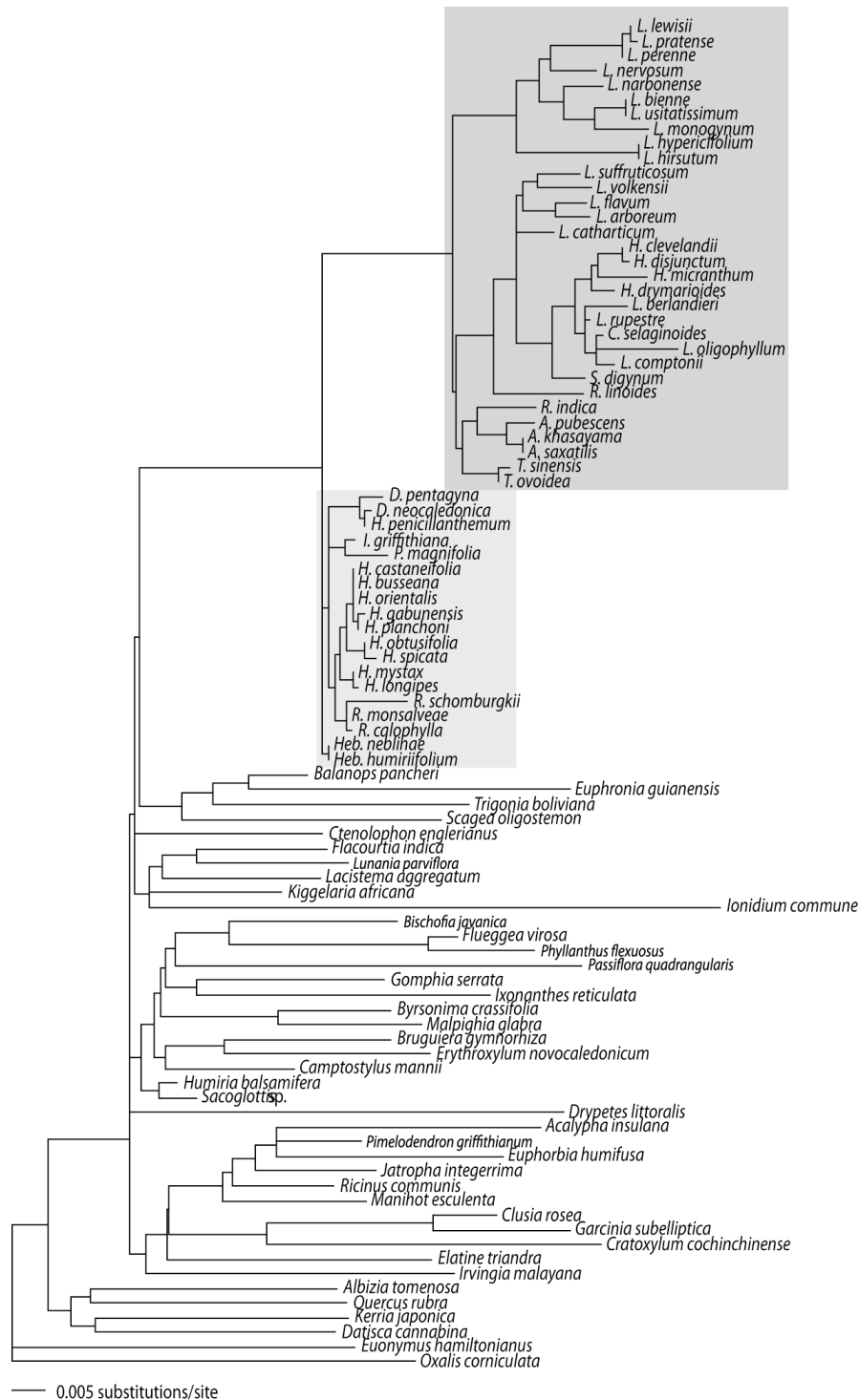


Fig. 2.5: ML phylogram for *rbcL*. Darker shading indicates Linoideae, lighter Hugonioideae.

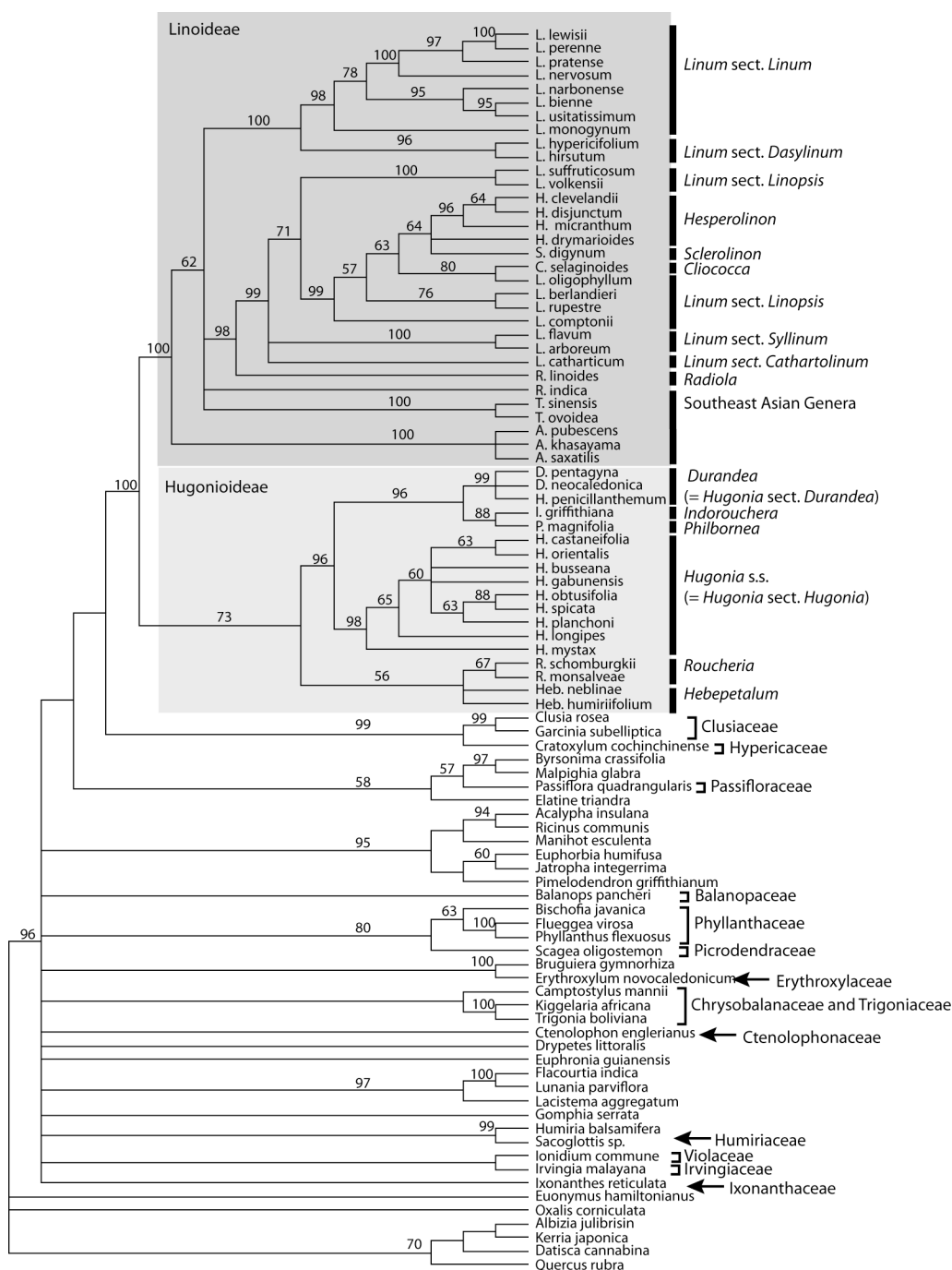


Fig. 2.6: Strict consensus of 1898 most parsimonious trees (L=3691) recovered during search on matK data alone, with MP bootstrap values above branches (values <50% not shown).

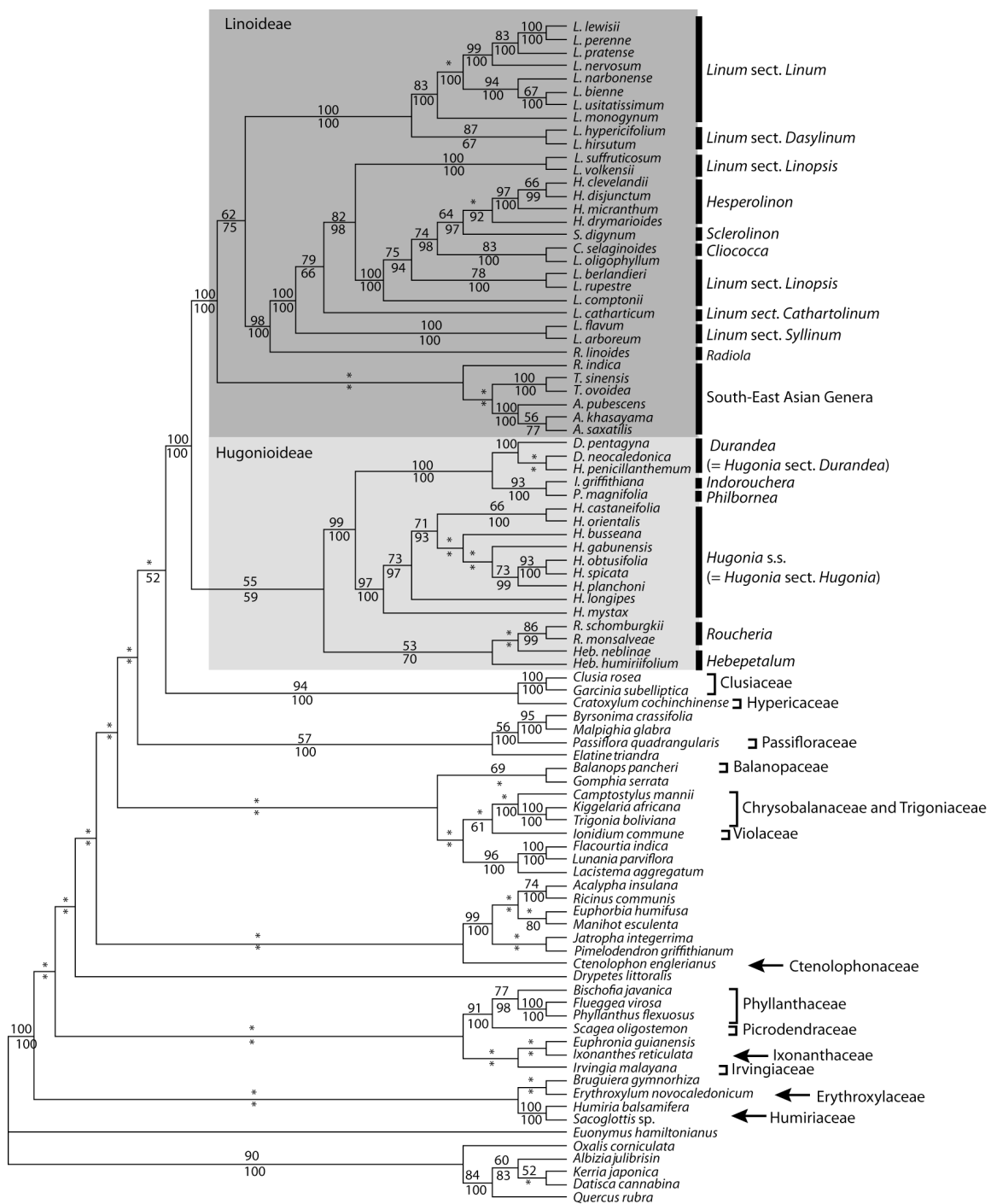


Fig. 2.7: ML topology for matK recovered by GARLI, with ML bootstrap values above branches, Bayesian posterior probabilities (as %) below (support values <50% indicated by \*).



Fig. 2.8: ML phylogram for matK data. Darker shading indicates Linoideae, lighter Hugonioideae.

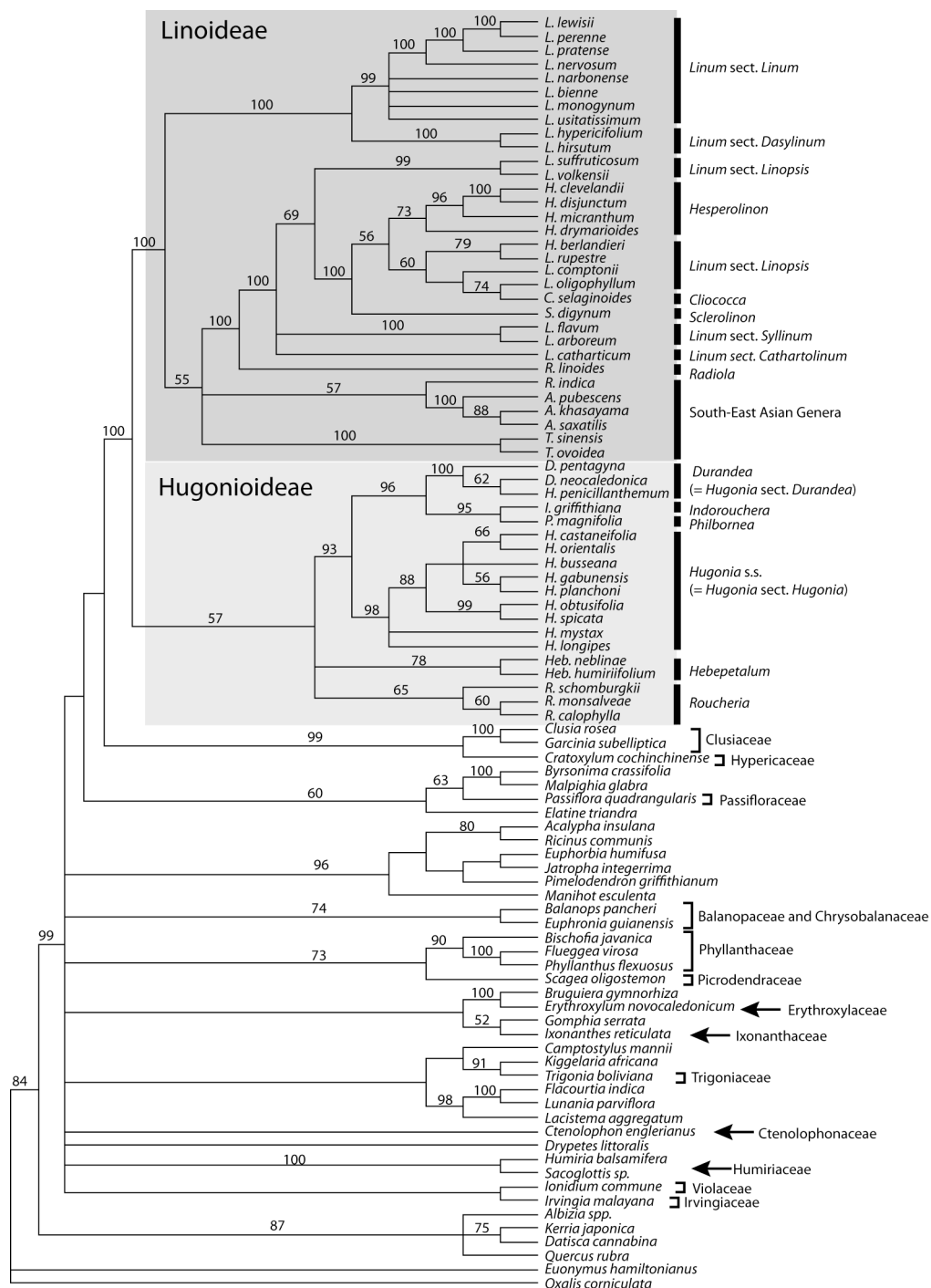


Fig. 2.9: Strict consensus of 314 most-parsimonious trees (L=5491) recovered from search on combined rbcL+matK matrix. MP bootstrap values above branches (values <50 not shown).

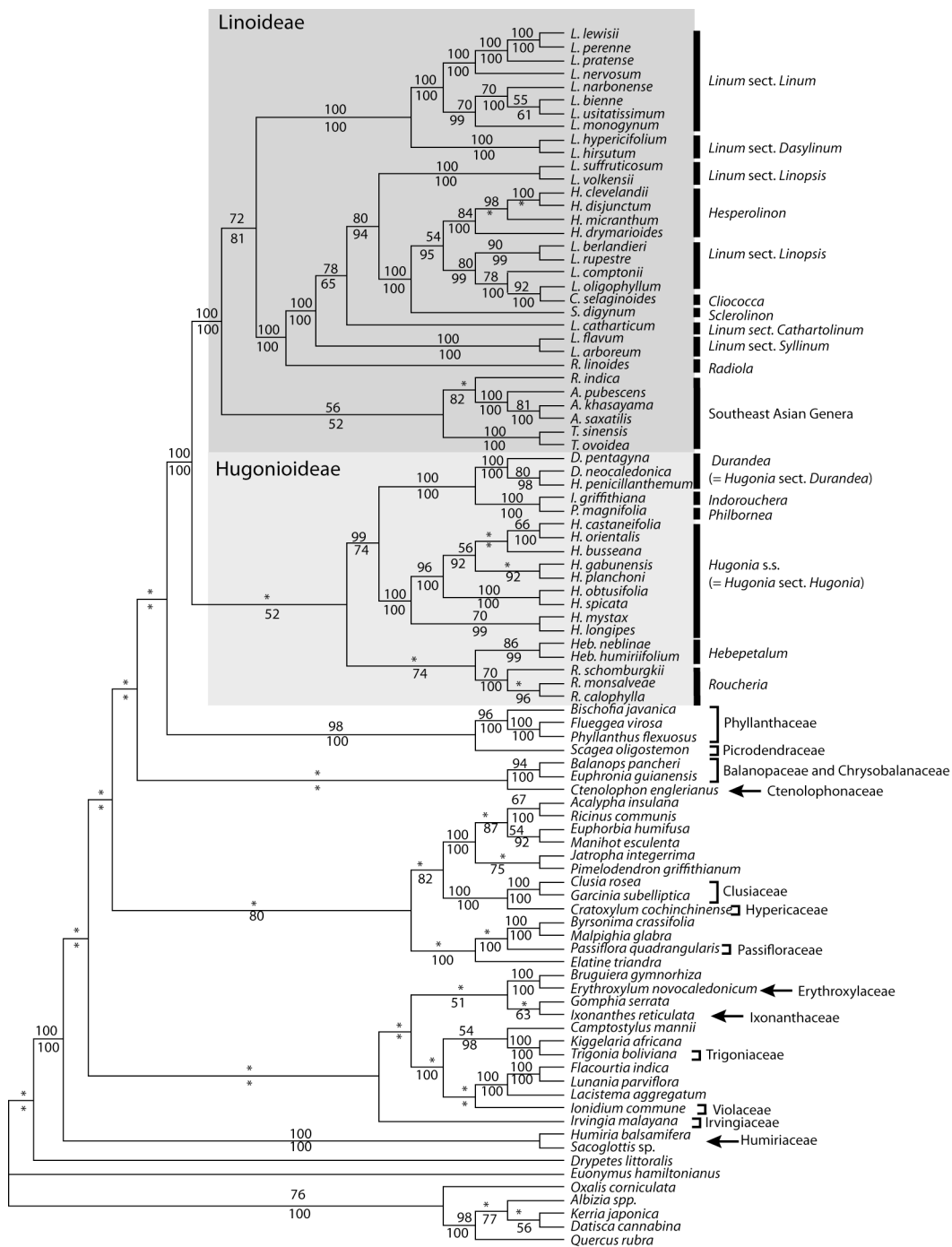


Fig. 2.10: ML topology for combined data recovered by RAXML using a partitioned model, with ML bootstrap percentages above branches and Bayesian posterior probabilities (as %) below branches. Values <50% indicated by \*.

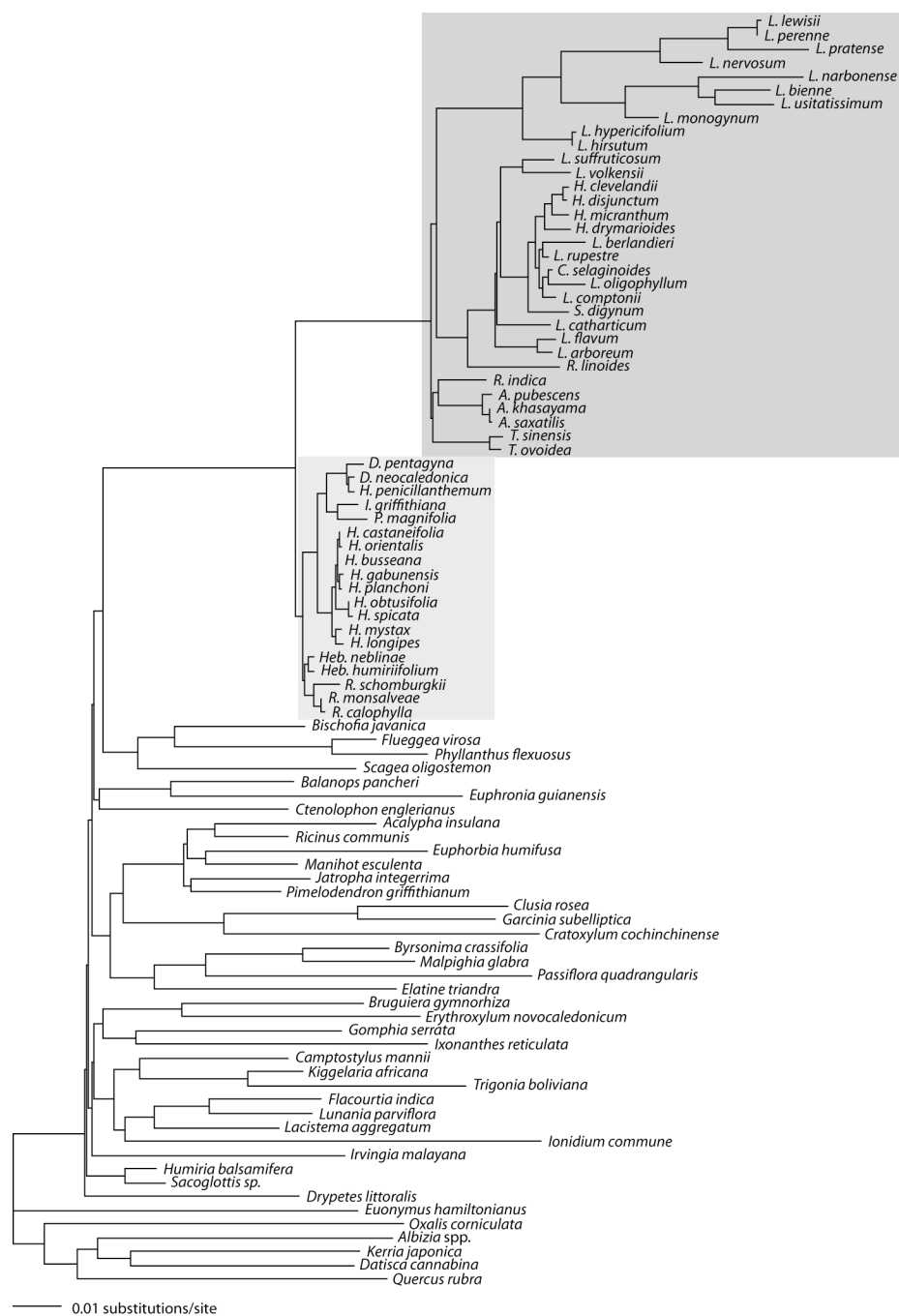


Fig. 2.11: ML phylogram recovered from combined data matrix by RAxML. Darker shading indicates Linoideae, lighter Hugonioideae.

Fig. 2.12: Chronogram (maximum clade credibility topology) resulting from BEAST analysis with the split of Linaceae and Irvingiaceae set at 105 mya, the maximum optimal divergence time estimated by Davis et al. (2005). Node ages set at average divergence time estimated by BEAST; grey bars indicate 95% height density proportion confidence intervals around nodes listed in Table 2.6. Clade posterior probabilities based on the post-burn-in tree sample indicated above branches (only values > 0.5 shown). Node D (common ancestor of all *Linum* and segregate genera) not shown due to non-monophyly of that grouping in the maximum clade credibility tree. Time scale given in millions of years before present. Major geologic time intervals are also indicated (pl = Pliocene, p = Pleistocene).





### **Chapter 3: Molecular Phylogenetics of of the Yellow-flowered flaxes, with Extended Sampling from *Linum* section *Linopsis* (Linaceae)**

#### **ABSTRACT**

The phylogenetic relationships of *Linum* sect. *Linopsis* sensu Rogers are examined using DNA sequences from 70 of the 94 species in the group, and 26 species representing the other four sections of *Linum* and the four currently-recognized segregate genera, *Cliococca*, *Hesperolinon*, *Radiola*, and *Sclerolinon*. Separate and combined analyses of the nuclear (ITS) and chloroplast (*matK*, *trnK* 3' intron, *ndhF*, *trnL-F* intron and spacer) data show that section *Linopsis* is paraphyletic given the current generic circumscriptions, and that its members are divided among five geographically distinct lineages which collectively are sister to *Radiola linoides*. The Northern Hemisphere Old World species of section *Linopsis*, along with sampled species of *Linum* sections *Cathartolinum* and *Syllinum*, appear either as a sister clade of, or basal paraphyletic grade to, a well-supported South African and New World lineage. The South African and South American species of section *Linopsis* are each well-supported monophyletic lineages, with the South American segregate *Cliococca* associated with South American *Linum*. The North American segregates *Hesperolinon* and *Sclerolinon* appear to be associated with some endemic Mexican species of *Linum*, while the remaining North American species comprise a large clade that may be sister to the South American lineage. Some previously circumscribed infrasectional taxa of section *Linopsis* are found to be well supported monophyletic groups: subsections *Dichrolinum*, *Halolinum*, and *Rigida*, and subsection *Linopsis* ser. *Virginiana*. We show that several of the characters generally used in *Linum* taxonomy are evolutionarily labile, but one feature, reduction of capsule dehiscence to produce a five-sectioned fruit, may be a unique morphological

synapomorphy for a lineage of North American flaxes (section *Linopsis* subsect. *Rigida* ser. *Rigida*).

## INTRODUCTION

In the genus *Linum* L., section *Linopsis* (Rchb.) Engelm. is the largest and most widespread of the five commonly recognized sections, with approximately 93 species distributed in Eurasia, Africa, and the Americas. The section includes annual and perennial species, most with pentamerous yellow flowers (pink or white in some species) with free petals and gland-fringed sepals (Rogers 1982). As in the other sections of *Linum*, the fruits of section *Linopsis* are dehiscent capsules composed of (usually) five carpels. Leaf arrangement, chromosome number, style and stigma morphology, capsule dehiscence pattern, pollen morphology, and the occurrence of stipular glands are features that vary among (and sometimes within) the species of section *Linopsis*.

Some sections of *Linum* are easily recognized by their distinctive morphological features. The species of section *Dasylinum* (Planch.) Juz., for instance, are known for their prominent pubescence, while those of section *Syllinum* Griseb. are distinguished by coherent petals and decurrent leaf bases. Section *Linopsis*, however, lacks any such obvious, diagnostic morphological characteristics. As a result, and due to the broad morphological variation and geographic distribution the section encompasses, the taxonomic concepts applied to these yellow-flowered flaxes have been particularly labile. Over the last 170 years, a shifting array of sections, subgenera, series, and segregate genera have been proposed in attempts to organize these species into a meaningful

hierarchy (Reichenbach 1837, Planchon 1847, 1848, Engelmann 1852, Bentham 1862, Gray 1865, Reiche 1894, Small 1907, Winkler 1931).

Most recently, Rogers and colleagues produced taxonomic revisions of the yellow-flowered flaxes of North America (Rogers 1963, 1968), South America (Mildner 1971, Mildner and Rogers 1978), and southern Africa (Rogers 1981a and 1981b). These works led to a revised infrasectional classification of species in section *Linopsis* (detailed in Rogers 1982 and 1984), built on the earlier systems of Planchon (1847, 1848) and Winkler (1931). Rogers' classification recognized five subsections and several sub-subsectional series within section *Linopsis*, with the characteristics and membership summarized in Table 3.1. Species sharing suites of similar characteristics were grouped together in his classification, yet few of the infrasectional taxa exhibit any unique, diagnostic features. Due to the continuous nature of variation in many characters, and the presence of species interpreted to be of “intermediate” or “transitional” morphology, Rogers maintained *Linopsis* as a section of *Linum*, preserved the most distinctive variants as valid segregate genera (*Cliococca* Bab., *Hesperolinon* (A. Gray) Small, *Sclerolinon* Rogers), and left *L. catharticum* L. as the sole species of *Linum* sect. *Cathartolinum* (Rchb.) Griseb. Some authors, including Small (1907), Winkler (1931), and Diederichsen and Richards (2003), have embraced more expansive concepts of *Cathartolinum* at various ranks, which included most of the North American species placed by Rogers in section *Linopsis*. Others have proposed additional North American yellow-flowered segregate genera (e.g. *Nezera* Raf., *Mesynium* Raf., *Mesyniopsis* W.A. Weber), but these have been only sporadically accepted (Löve 1982, Weber 1984).

Based on the assumption that the Mediterranean region was the “ancestral area” for *Linum*, and that species found there would thus exhibit “ancestral” character states, Rogers proposed an hypothesis of evolutionary relationships in section *Linopsis* (Rogers 1982 Fig. 1, redrawn here in Fig. 3.1). The subsections of section *Linopsis* were depicted as originating from an Old World ancestor (the “ancestral type”), presumed to be a perennial, heterostylous species with free styles, incomplete septa, and tricolpate pollen. Rogers’ subsection *Linopsis* includes species with this same set of ancestral features (except heterostyly, which Rogers assumed had been evolutionarily “lost” in most species). The other subsections and series of section *Linopsis* were recognized based on the occurrence of “derived” character states, such as fused styles (as in the single species of series *Mysorensis*), unusual chromosome numbers ( $n=13$  in the three species of series *Neo-Mexicana*, for instance), and distinctive corolla pigmentation (as in the white or pinkish flowers of subsection *Dichrolium*). The New World subsection *Rigida*, with fused styles, multiporate pollen, and, generally, complete septa, was considered the most “advanced” subsection, exhibiting the greatest number of “derived” character states. Rogers’ classification and depiction of relationships within the section thus predict that neither subsection nor series *Linopsis* are monophyletic, by showing other subsections and series arising from different points within them. Subsections or series that are characterized by derived states, such as subsection *Rigida*, might be expected to be monophyletic lineages.

The unique morphology of the South American segregate *Cliococca* left its evolutionary origins mysterious (Rogers and Mildner 1971) until McDill et al. (2009)

showed its single species, *C. selaginoides* (Lam.) C. M. Rogers, to be sister to the four South American species of *Linum* sect. *Linopsis* sampled in their analyses. While several features of *Cliococca*, such as its rhizomatous habit, are not found in *Linum*, it is reported to share the same chromosome number ( $n=18$ ) reported from South American *Linum* (Rogers and Mildner 1971) and from many North American species (Rogers 1984).

McDill et al. (2009) also found the North American segregates, *Hesperolinon* and *Sclerolinon*, to be sister to each other, and closely related to New World *Linum* sect. *Linopsis*. The similarities and possible close relationship of *Hesperolinon* and *Sclerolinon* were noted by Gray, who placed these species together in *Linum* sect. *Hesperolinon* (Gray 1865). Both genera exhibit a reduction in carpel number relative to five-carpellate *Linum*, and are annual species of western North America (whereas most North American *Linum* are perennials occurring east of the Rocky Mountains). Rogers (1975) hypothesized the relationship of *Hesperolinon* to the North American series *Neo-Mexicana* based on pollen and floral morphology. Xavier (1967) also noted similarities in the pollen morphology of *Hesperolinon*, *Sclerolinon*, and the members of series *Neo-Mexicana*. In their preliminary investigation of the *Linum* phylogeny, McDill et al. (2009) included one member of series *Neo-Mexicana*, *L. kingii*, which did not appear as closely related to *Hesperolinon* or *Sclerolinon*. The origins of these segregates may lie among the previously unsampled species North American *Linum* sect. *Linopsis*. Of particular interest in this matter may be certain endemic Mexican species of *Linum*, such as *L. schiedeana* Schldl & Cham. and *L. mexicanum* Kunth, with basal whorled leaves

that recall those observed in some *Hesperolinon*. Most other North American species of *Linum* have entirely alternate leaves (Rogers 1975).

The investigation of the molecular phylogeny of the temperate Linaceae by McDill et al. (2009) revealed that members of *Linum* sect. *Linopsis* are intermingled with those of two Old World sections of *Linum* (*Cathartolinum* and *Syllinum*), as well as the three segregate genera. By finding *Linum* and its largest section to be non-monophyletic, McDill et al. (2009) revealed the potential need for a revised taxonomy of *Linum* and its sections and segregates, but that study left many questions unanswered due to limited taxon sampling. Their analyses showed that New World and South African linums are probably derived from Eurasian ancestry, supporting Rogers basic assumptions of the origin of section *Linopsis* and potentially his polarization of character evolution. However, since the vast majority of New World and African taxa were not sampled, it remains to be seen whether any of Rogers' subsectional taxa, or any alternative groupings proposed by previous workers for species of section *Linopsis*, are monophyletic.

In this study, using greatly expanded sampling of African and American species of *Linum* sect. *Linopsis*, we further explore the relationships among sections *Linopsis*, *Cathartolinum*, and *Syllinum*, and the three segregate genera and assess the monophyly of the subsections and series of section *Linopsis* proposed by Rogers (1982, 1984). We also examine the evolution of several morphological characters used by Rogers and others as indicators of relationships to determine character polarity and evaluate the utility of these characters as a basis for classification or as synapomorphies of various lineages. Finally,

our phylogenetic hypotheses will provide a context for testing theories about the biogeographic history of this diverse, wide-spread lineage (Chapter 5).

## **MATERIALS AND METHODS**

**Taxon Sampling** —As a basis for sampling from *Linum* sect. *Linopsis*, we accepted the species and taxonomic groupings recognized by C.M. Rogers and R. Mildner in their treatments of *Linum* and its segregate genera in North America (Rogers 1984), South America (Rogers and Mildner 1971, Mildner and Rogers 1978), and Africa (Rogers 1981a, 1981b). Seventy (indicated by asterisks in Table 3.1) of the approximately 94 species of section *Linopsis* (including one undescribed species) are represented in our analyses. Members of *Linum* sects. *Cathartolinum* (1 species sampled out of 1 currently recognized) and *Syllinum* (5/37) were selected for inclusion, along with representative samples of the segregate genera *Cliococca* (1/1), *Hesperolinon* (4/13), *Sclerolinon* (1/1), and *Radiola* (1/1), all of which were found to be associated with section *Linopsis* by McDill et al. (2009). Representative species from the blue-flowered sections of *Linum* (12/63) and the southeast Asian genus *Reinwardtia* (1/2) are used as outgroups for the chloroplast (CPL) data analyses. The 136 DNA accessions, representing 95 different species of yellow-flowered *Linum*, segregate genera, and outgroups included in this study are listed in Appendix B along with voucher specimen information for each accession and morphological characters states scored for each species.



**DNA Extraction, PCR, and Sequencing** — DNA extractions utilized leaves either collected fresh and silica-dried, or judiciously garnered with permission from existing herbarium specimens. Extractions were performed using the modified Doyle and Doyle CTAB protocol (Loockerman and Jansen 1996). Between 2 and 6 microliters of a 1:10 dilution (in TE) of each extract were used as template in PCR reactions. Sequences from four regions of the chloroplast genome were utilized in this study: a portion of the *ndhF* gene, the *trnL-F* intron and spacer, the *trnK* 3' intron, a portion of the *matK* gene. From the nuclear genome, the ribosomal ITS was sequenced. Primers used for amplification and sequencing of the *trnL-F* regions, *trnK* 3' intron, and *ndhF* are as reported in McDill et al. (2009). Primers used to amplify *matK* for this study are provided in Table 3.2. ITS primers used are given in Simpson et al. (2004). For each PCR, a “master mix” was assembled that contained (per reaction): 2.5µL 10X Triton-X reaction buffer, 200 µM dNTPs, 10pM each primer, 2-3 µL 25 mM MgCl<sub>2</sub>, and *Taq* polymerase, in a total volume of 25µL. As in McDill et al. (2009), BSA (3µL of 3.3% w/v per reaction, Savolainen et al. 2000) was used to enhance or obtain amplification of some regions from some samples as necessary. The thermal cycling program for PCR consisted of the following steps: initial denaturation 95°C/4 min followed by 34 cycles with 95°C/30 sec denaturation , 50°C/30 sec annealing , and 72°C/30 sec extension . A final extension step of 6 min at 72°C was included to finish synthesizing any incomplete products. PCR products were visualized by electrophoresis on 1.5% agarose gels. An EXO-SAP protocol based on Werle et al. (1994) was used to eliminate unincorporated primers and nucleotides prior to sequencing. PCR products were sequenced with forward

and reverse dye terminator reactions using the amplification primers for each marker (internal primers were also employed for sequencing of *matK*), at The University of Texas at Austin's Institute for Cell and Molecular Biology Core Facility.

Sequencher 4.1.2 or 4.5 (GeneCodes Corp. 2005) were used to assemble and edit forward and reverse sequence reads. Alignments of *matK*, *ndhF*, and the *trnK* 3' intron were produced manually in MacClade 4.08 (Maddison and Maddison 2005), while *trnL-F* and ITS sequences were aligned using MUSCLE (Edgar 2004), followed by manual adjustment in MacClade. Areas of uncertain or ambiguous alignment were excluded from analyses; gaps were treated as missing data.

**Phylogenetic Analyses** —Prior to analysis, an appropriate substitution model was determined for each marker using the Akaike Information Criteria (AIC) implemented in Modeltest 3.7 (Posada and Crandall 1998), so that the appropriate model types could be applied during the analyses. Maximum likelihood (ML) and Bayesian analyses were subsequently performed on three data matrices: the combined chloroplast (CPL) matrix, the ITS matrix, and the combined CPL+ITS matrix.

Maximum-likelihood analyses were conducted with GARLI v. 0.96b (Zwickl 2006), with the appropriate substitution model type (as determined by ModelTest) implemented but specific parameter values left to be optimized by GARLI. Since this version of GARLI does not allow data partitioning, the model used for the CPL and combined CPL+ITS analyses applied a single GTR+G+I model to the entire matrix. All settings in GARLI were left at their default values. One-hundred GARLI search replicates were executed for each matrix, to assess the occurrence of local optima. Non-

parametric ML bootstrap values for each matrix were determined from 500 bootstrap replicates performed with GARLI, with one search replicate per bootstrap replicate.

For Bayesian analyses, MrBayes v.3.1.2 (Huelsenbeck and Ronquist 2001) was used. Data matrices consisting of combined genomic regions (i.e., the CPL matrix and the CPL+ITS matrix) were partitioned, and the appropriate form of substitution model (but not the specific estimated model parameter values themselves) were applied for each data subset (region). Prior probability distributions on all parameters were kept at default values, and all parameters except for tree topology and branch lengths were unlinked between partitions. Two independent analyses were run simultaneously (nruns=2), with the built-in convergence diagnostic (average standard deviation of split frequencies) assuming a relative burn-in of 25%. Analyses were run for a minimum of 4 million MCMC generations (sampling rate = 0.01), at which point they were stopped if the convergence diagnostic had dropped to 0.01 or lower. If the two runs had not converged, the analysis was continued in two-million-generation increments until convergence to the desired level was obtained.

**Analysis of ITS** — Due to the potential for sequencing paralogous and pseudogenic copies of the nuclear ribosomal repeat (Feliner and Rosselló 2007), ITS PCR products for most samples were cloned using TOPO-TA cloning kits (Invitrogen). Multiple clones from each cloned sample were sequenced when possible (using the M13F and R vector primers included with the kit). Sequences exhibiting one or more changes within certain regions of the 5.8S ribosomal DNA known to be stringently conserved in angiosperms (Harpke and Paterson 2008) were designated as pseudogenes. A

preliminary ML phylogenetic analysis of all ITS sequences (using GARLI as described above) was performed to identify putative paralogous sequences, designated as any non-pseudogene ITS sequences from a single DNA sample which appear in different, well-supported clades in the ML bootstrap tree). A reduced ITS matrix was produced for subsequent ML and Bayesian analysis (as described above) by removing pseudogene and extraneous clone sequences. For samples with ITS clones determined not to represent multiple paralogous copies, a single clone sequence was chosen for inclusion in subsequent analyses. Any DNA sample that exhibited apparent paralogous sequences was represented in the subsequent ML and Bayesian analyses of the reduced ITS matrix by sequences of each ITS “type” found. In the combined CPL+ITS analysis, samples that exhibited putative paralogous ITS sequences are represented by their chloroplast marker sequences alone (with ITS coded as missing data).

**Analysis of Character Polarity and Evolution** — To examine character evolution and assess whether any morphological character states might characterize molecular lineages revealed by the foregoing analyses, states for selected characters used in Rogers’ classification were mapped onto the best tree topology resulting from the combined data ML analysis using MPRs (Most Parsimonious Reconstructions) mode in Mesquite v. 2.7 (Maddison and Maddison 2009). Character states scored for each species are provided in Appendix A.

## RESULTS

Data matrix characteristics and statistics for each marker are provided in Table 3.3.

**Missing Data** — The following sequences were not obtained due to chronic amplification failures [note: *Linum* species listed here are members of section *Linopsis* unless otherwise noted]. Numbers associated with species names are lab accession numbers for DNA extracts, and correspond to voucher specimens listed in Appendix A. **matK:** *L. heterostylum* 62, *L. thunbergii* 067, *L. rzedowskii* 165 (section *Linum*). **trnK 3' intron:** *L. schiedeanum* 9, *C. selaginoides* 49, 206, 220, *L. littorale* 89, *L. oligophyllum* 91, LPB2, *L. cremnophyllum* 187, *L. brevifolium* 193, *L. prostratum* 194, *L. chamissonis* 195, *L. erigeroides* 196, *L. carneum* 200, *L. macraei* 210, *L. chamissonis* 215, *L. scoparium* 219, *L. burkartii* 221, *L. hudsonioides* 244, *L. carteri* 266, *L. aethiopicum* SA12. **ndhF:** *L. quadrifolium* 258. **trnL-F intron and spacer:** *L. berlandieri* 27, *L. hudsonioides* 032, 244, *L. erigeroides* 196, *L. intercursum* 401. **ITS:** *L. oligophyllum* LPB2, *L. rigidum* 400, *L. berlandieri* A1, *L. subteres* 302, *L. berlandieri* 241, *L. aristatum* 233; no ITS sequences were included from blue-flowered linums or *Reinwardtia indica*. Sequence alignments for the chloroplast and ITS data as analyzed in these analyses are presented in Appendices C and D, respectively.

**Chloroplast Phylogeny** — The phylogeny of combined chloroplast marker sequences (136 individuals representing 95 species, including 12 blue-flowered linums and the rooting outgroup *Reinwardtia indica* ) are shown in Fig. 3.2 (ML phylogram) and

Figs. 3.3 and 3.4 (Bayesian majority-rule consensus with ML bootstrap (BS) and posterior probability (PP) values).

The blue-flowered linums are found in a well-supported monophyletic group, with the two members of *Linum* sect. *Dasylinum* sister to sect. *Linum* (Fig. 3.3). Sampled species of yellow-flowered *Linum* and the segregate genera are shown to form a well-supported monophyletic group sister to *Radiola* (Fig. 3.2). Several well-supported lineages of Old World yellow-flowered linums are evident, but the branching order among them is not well supported (Figs. 3.3, 3.4). *Linum* sect. *Syllinum* is shown as monophyletic, but section *Linopsis* is not. Section *Linopsis* subsect. *Halolinum* appears as sister to section *Syllinum*. *Linopsis* subsect. *Dichrolinum* is shown as sister to the Old World Northern Hemisphere (OWNH) members of subsection *Linopsis* ser. *Linopsis* (Figs. 3.2, 3.3). The position of *Linum* sect. *Cathartolinum* is not well supported. In the ML topology, *L. catharticum* appears sister to a clade comprised of *Linum* sect. *Linopsis* (excluding subsection *Halolinum*) and the segregate genera, but that relationship does not obtain significant bootstrap support (Fig. 3.2). In the Bayesian consensus topology (Fig. 3.3), *L. catharticum* appears as sister to the South African and New World clade of section *Linopsis* and the segregate genera.

The lineage containing the South African and New World Yellow flowered linums is well supported, with the New World group sister to the clade of South African species (Fig. 3.3). Relationships within the South African clade are not well resolved, although some groupings are well supported.

The New World taxa of section *Linopsis* and the segregate genera are divided among four moderately-to-well supported clades (Fig 3.4): (1) two species (*L. kingii* and *L. neomexicanum*) of subsect. *Linopsis* ser. *Neo-Mexicana*, (2) the six sampled species of subsect. *Linopsis* ser. *Virginiana*, (3) a clade including subsect. *Rigida*, *L. pringlei* ( of subsection *Linopsis* ser. *Neo-Mexicana*), and some North American members of subsect. *Linopsis* ser. *Linopsis*, and (4) a clade including the remainder of North and South American subsect. *Linopsis* ser. *Linopsis* and the segregate genera *Cliococca*, *Hesperolinon*, and *Sclerolinon*. Relationships among these four New World lineages are not resolved with significant levels of bootstrap or posterior-probability support.

Subsection *Rigida* is a well-supported lineage, as is the genus *Hesperolinon*, which is shown as sister to *Sclerolinon*. The South American taxa are a strongly supported lineage, with *Cliococca selaginoides* sister to South American linums. Many of the species of North American subsection *Linopsis* ser. *Linopsis* are divided among three well-supported clades, not recognized formally in Rogers' classification, but henceforth referred to as the *Linum rupestre* group (clade B in Fig. 3.4, consisting of *L. arenicola*, *L. flagellare*, *L. rupestre*, and *L. bahamense*), the *Linum schiedeanum* group (clade A in Fig. 3.4, *L. schiedeanum*, *L. cruciata*, *L. lasiocarpum*, *L. tenellum*, and *L. nelsonii*), and the *Linum mexicanum* group (clade C in Fig. 3.4, *L. mexicanum*, *L. guatemalense*, and *L. orizabae*). The *Linum rupestre* and *Linum schiedeanum* groups both appear related to subsection *Rigida*, while the *Linum mexicanum* group is associated with the South American linums and the segregate genera.

**ITS Phylogeny**— In the analysis of ITS clones, 460 cloned and direct sequences were included for the sampled taxa of the yellow-flowered lineage (Figs. 3.5-3.8). Only 36 sequences were definitively identified as pseudogenes based on the criteria of Harpke and Paterson (2008) and subsequently excluded. Cloned sequences from most samples formed exclusive, well-supported clades, from each of which a single non-pseudogene copy was selected for inclusion in further analyses. Six species exhibited apparent paralogous copies found in different moderately or well-supported clades: *L. adustum* SA8, *L. gracile* SA5, *L. westii* 269, *L. mexicanum* 37, *L. nelsonii* F5, and *L. guatemalense* (the first two species listed are South African, the rest North American).

Due to the variable nature of the ITS, and resulting difficulty of aligning sequences from yellow-flowered taxa with blue-flowered *linums* and *Reinwardtia*, the ITS-inclusive analyses utilize sequences from members of the yellow-flowered clade only, with *Radiola* used as the outgroup for tree rooting purposes. *Radiola linoides* was found as sister to the yellow-flowered clade by McDill et al. (2009), and in the CPL phylogeny presented here (Fig 3.3).

Results from the ML and Bayesian analyses of the reduced ITS matrix (Figs. 3.9 and 3.10, respectively) reveal many of the same well-supported lineages as the CPL analyses, but also lack strongly-supported resolution of relationships among those lineages. There are however several notable divergences from the CPL phylogeny. The position of section *Cathartolinum* (sister to section *Linopsis* subsect. *Halolinum*, Fig. 3.10) differs from that found in the CPL analyses (where it was sister to the New World



and South African clade. In the ITS topology, all of the Old World Northern Hemisphere yellow-flowered linums comprise a moderately-supported clade sister to the African-New World clade. The relationship of South African linums to the New World taxa is not well resolved in the ITS analysis. The *Linum schiedeanum* group is not monophyletic in the ITS analyses: two of its species are a sister clade to *Linopsis* ser. *Virginiana* (PP=96%, BS <50%), while the remainder form a clade (PP=100%, BS<50%) whose position within the New World lineage is unresolved (Fig. 3.10, group A, the *L. schiedeanum* group, shown in two parts). Section *Linopsis* subsect. *Rigida*, monophyletic in the CPL results, is non-monophyletic in the ITS topologies (Fig. 3.10) due to the placement of *Linum pringlei* (of subsection *Linopsis*) as sister to *L. sulcatum* (86% PP, 89% BS).

Putative paralogous sequences from the South African species *Linum adustum* (indicated by black stars in Figs. 3.9 and 3.10) and *L. gracile* (black dots) are dispersed among two clades, one with sequences from two other accessions of *L. gracile* plus *L. heterostylum*, and the other consisting of sequences from *L. aethiopicum*. Sequences isolated from the Mexican species *Linum nelsonii* (triangles, Fig. 3.10) appear variously related to those from two other Mexican taxa, *L. schiedeanum* and *L. cruciata*. Putative paralogs from *L. mexicanum* (squares) and *L. guatemalense* (diamonds) are confined within the *Linum mexicanum* group of subsection *Linopsis* (group C, Figs. 3.9 and 3.10), which is shown as sister to the *Hesperolinon*+*Sclerolinon* lineage (98% PP, <50% BS). Paralogs from *L. westii* (asterisks in Fig. 3.10) are widely separated in the phylogeny: two fall among species of series *Virginiana* (which agrees with the chloroplast marker

placement of *L. westii*, and the classification of Rogers), while the third is nested within subsection *Rigida* (sister to a sequence from *Linum aristatum*).

**Combined Data Phylogeny** — ML and Bayesian analyses of the combined CPL+ITS data matrix (paralogous ITS sequences excluded) are presented in Figs. 3.11 and 3.12, respectively. These results mimic those from the individual CPL and ITS analyses described above, featuring poorly supported relationships within and among a few well-supported lineages. *Linum catharticum* is moderately supported as sister to a clade containing the other Old World Northern Hemisphere species. The South African species of *Linopsis* form a well-supported clade, again sister to the New World taxa. A clade containing *Hesperolinon*, *Sclerolinon*, and the *Linum mexicanum* group of subsection *Linopsis* ser. *Linopsis* receives tenuous support as the most basal New World lineage. The clade of South American *Linopsis*, *Cliococca*, and the Mexican species *L. longipes*, is very weakly supported as monophyletic and sister to a large North American clade. Within that group, the South American taxa are well supported as monophyletic, with *Cliococca* nested among the *linums* (sister to *L. burkartii*, as in the ITS analyses). Series *Virginiana* is well supported, as is series *Neo-Mexicana* (excluding *L. pringlei*, which appears as sister to the *Linum rupestre* group).

**Character Mapping** — Fruit segmentation, pollen morphology, extent of false septa, and stylar fusion are mapped on the combined data ML topology in Fig. 3.13.

## DISCUSSION

While the relationships among the lineages of yellow-flowered linums, particularly among the New World species, remain somewhat poorly resolved or poorly supported, several major conclusions are possible.

**Relationships among the “yellow-flowered” sections of *Linum*** — Three sections of *Linum* fall into what has been called the yellow-flowered clade (McDill et al. 2009): *Linopsis*, *Syllinum*, and *Cathartolinum*. Section *Linopsis* is clearly not monophyletic as circumscribed by Rogers (1982), with *Linum* sects. *Syllinum* and *Cathartolinum*, as well as the segregate genera *Cliococca*, *Hesperolinon*, and *Sclerolinon* nested within it. This reiterates the findings of McDill et al. (2009), but with far more extensive taxonomic sampling from section *Linopsis*. *Syllinum*, a Eurasian section well-defined by morphological features including coherent petals and decurrent leaves, is consistently found to be monophyletic and sister to the Mediterranean subsection *Halolinum* of section *Linopsis*, as found by McDill et al. (2009). The relationships of section *Cathartolinum*, with its single species, *Linum catharticum*, are unclear. As found by McDill et al. (2009), chloroplast and nuclear markers appear to provide conflicting signals, with the CPL indicating *L. catharticum* as sister to the African+New World lineage, and the ITS showing *L. catharticum* nested among Eurasian taxa. While the placement of *L. catharticum* in the ITS topology, nested within the Old World Northern Hemisphere (OWNH) clade, is moderately-to-well-supported by bootstrap values and posterior probabilities (Fig. 3.10), the branching order among the Old World Northern Hemisphere lineages in the CPL phylogeny (Fig. 3.3) is not well supported. In the

combined analysis, *L. catharticum* assumes a “compromise” position, sister to an Old World clade. The possible taxonomic implications of the placement of *L. catharticum* are discussed more extensively by McDill et al. (2009).

**Relationships among the members of *Linum* section *Linopsis*** — As noted above, section *Linopsis* as circumscribed by Rogers is not monophyletic, and its species are divided among several major lineages. Some of these lineages correspond to Rogers’ subsections or series. Among the Old World lineages of *Linum*, subsections *Halolinum* and *Dichrolinum* appear to be well-supported groups in the molecular phylogenies, and are well-defined by other characters: *Halolinum* with a unique base chromosome number ( $x=10$ ), and *Dichrolinum* with white or purplish flowers (Rogers 1982).

In the New World, subsection *Rigida* also appears to be a well-defined and supported lineage, exhibiting several morphological features (such as united styles, multiporate pollen, 5-segmented capsules, and  $n=15$ ) designated by Rogers as “derived” states (Rogers 1982). *Linum sulcatum* (subsection *Rigida* ser. *Sulcata*), was considered by Rogers (1982) to be “intermediate” between subsection *Rigida* ser. *Rigida* and members of subsection *Linopsis* due to its united styles (a feature of series *Rigida*) and 10-segmented capsules (a feature of subsection *Linopsis*). In the combined data analyses, *Linum sulcatum* appears as sister to series *Rigida*, in what could be considered an “intermediate” position between that series and the members of subsection *Linopsis* (Fig. 3.12), as Rogers might have predicted (the ITS data indicate a different, anomalous placement of *L. sulcatum*, sister to *L. pringlei*, with moderate support (pp = 86%, ML bootstrap 89%).

Series *Virginiana* (of subsection *Linopsis*) is also consistently found to be monophyletic and well-supported, with six of its seven species sampled. This group of herbaceous perennials is confined to eastern North America. Some of the species of series *Virginiana* exhibit derived characters, such as complete septa and colporate pollen.

Rogers' subsection *Linopsis* is not monophyletic, as subsections *Dichrolinum*, *Halolinum*, and *Rigida* are nested within it. The non-monophyly of section and subsection *Linopsis* is not surprising, since species were essentially assigned to these groups based their *lack* of derived features, rather than the presence of synapomorphies. The limited power of plesiomorphic character states to predict relationships, and their propensity to define paraphyletic groups when used for classification purposes, are well known (e.g. Donoghue and Cantino 1988). That said, the non-monophyly of section and subsection *Linopsis* was also essentially predicted by C.M. Rogers; the evolutionary scenario he proposed indicates a nested series of relationships with different subsections and series arising from different points within subsection *Linopsis* (see Fig. 3.1 and Rogers 1982, Rogers 1969). The classification of *Linum* section *Linopsis* proposed by Rogers (Table 3.1) was thus not designed to represent hypothetical monophyletic lineages as infrasectional taxa, but rather as a way of indicating morphological similarity. Species were organized into subsections and series based on shared suites of features, and evolutionary relationships described verbally or graphically (Rogers 1969, 1982), with more “advanced” groups arising from within paraphyletic ancestral taxa.

Our analyses suggest that several species of subsection *Linopsis* ser. *Linopsis* are members of three lineages that have not been formally recognized in classifications: the

“*Linum schiedeanum* group” (*L. schiedeanum*, *L. cruciata*, *L. lasiocarpum*, *L. tenellum*, and *L. nelsonii*), the “*Linum rupestre* group” (*L. rupestre*, *L. bahamense*, *L. flagellare*, and *L. arenicola*), and the “*Linum mexicanum* group” (*L. mexicanum*, *L. guatemalense*, *L. orizabae*).

The *Linum schiedeanum* group is well supported by chloroplast data and combined analysis (group A, Figs. 3.4, 3.12), but not the ITS alone (Fig. 3.10). This group is characterized by whorled leaves, an unique feature among North American linums but also found in some South American and South African species. The species of the *Linum schiedeanum* group are distributed from the Guadalupe mountains of Western Texas south to Veracruz.

The *Linum rupestre* group (clade B, Figs. 3.4, 3.9, 3.10, 3.11, 3.12) is characterized by a perennial habit with many branches originating from a woody taproot or caudex, and most of the species occur in calcareous substrates (Rogers 1984). *L. rupestre* is found on limestone outcrops from central Texas to central Mexico. *Linum bahamense* and *L. arenicola* are found in similarly calcareous soils in the Bahamas and Florida, respectively. *Linum flagellare*, however, is occurs in sandy soils in Coahuila, Zacatecas, and Durango (Mexico).

The three species of the *Linum mexicanum* group (clade C, Figs. 3.4, 3.9, 3.10, 3.11, 3.12) are native to central and southern Mexico. Similarities among the three species of this group were noted by Rogers (1984), who characterizes them as distinctive “coarse” perennial herbs or subshrubs. Leaves in these species are opposite or whorled at the base of the plant, transitioning to alternate toward the inflorescence. The leaf

arrangement transition from whorled or opposite at the base to alternate in the upper portions of the plant is also observed in some *Hesperolinon* and in *Sclerolinon*, which may share common ancestry with the *L. mexicanum* group (Fig. 3.12).

An anomalous result that contradicts Rogers classification in spite of the presence of a “derived” feature is the non-monophyly of series *Neo-Mexicana* (of subsection *Linopsis*), whose three species are reportedly characterized by a unique base chromosome number of  $x=13$  (Rogers 1984). *Linum kingii* and *L. neomexicanum* (both  $n=13$ ) are known from high elevation mountain ranges and “sky islands” from Colorado to Nevada and south to New Mexico, while the tetraploid *L. pringlei* ( $n=26$ ) occurs in the Sierra Madre of the Mexican states Sonora and Chihuahua. *Linum kingii* and *L. neomexicanum* appear to be sister species our results, but the affinities of *L. pringlei* are unclear. The chloroplast phylogeny (Fig. 3.2) and combined analysis (Fig. 3.12) show *L. pringlei* sister to the *L. rupestre* group of species, while ITS places it with subsection *Rigida*.

Unfortunately *L. pringlei* ITS was not cloned, so we do not know whether the direct sequence represents a paralogous sequence copy, which could explain the disagreement between the ITS and CPL placements for this taxon. It may also be that the  $n=26$  chromosome number of *L. pringlei* is not a tetraploid derivative from the  $n=13$  of *L. kingii* and *L. neomexicanum* as Rogers thought, but is derived from some other ancestral number via some combination of polyploidy and aneuploid change.

**Relationships of *Cliococca*, *Hesperolinon*, and *Sclerolinon*** — The expanded sampling from *Linum* sect. *Linopsis* provides a more detailed perspective on the origins and relationships of the three segregate genera and raises some additional questions.

McDill et al. (2009) found the South American segregate *Cliococca* sister to the four species of South American *Linum* sampled in their study. While the chloroplast phylogeny presented here gives the same result (*Cliococca* sister to South American *Linum*, PP=98 BS=66, Fig. 3.4), the ITS and combined analysis show *C. selaginoides* derived from within the South American *Linum* clade, and perhaps sister to *L. burkartii* (Figs. 3.10, 3.12). Though the ITS from all three accessions of *C. selaginoides* were cloned, and those clones consistently formed a monophyletic grouping sister to a clade of *L. burkartii* clones (Fig. 3.6), most South American taxa were not cloned, so the possibility of non-homologous ITS copies affecting the result can't be eliminated. While *Cliococca* is reported to share the same chromosome number as South American *Linum* species ( $n=18$ ), its divergent morphology yet makes it difficult to recognize any synapomorphies it might share with other species (Rogers and Mildner 1971). Its close relationship to South American linums, though, is apparent.

As in McDill et al. (2009), the North American segregates *Hesperolinon* and *Sclerolinon* are shown to be sister to each other. Species of these genera were placed by Gray in his *Linum* sect. *Hesperolinon*, and are characterized by a reduction in carpel number from the usual five found in *Linum*, a tendency toward opposite or whorled leaves, and distributions in Western North America (most North American species of *Linum* occur east of the Rocky Mountains). Small (1907) segregated the genus *Hesperolinon*, but did not include *L. digynum*, instead placing it in the genus *Cathartolinum* along with the other North American yellow-flowered species (Small 1907). In her seminal revision of *Hesperolinon*, Sharsmith (1961) described several new



*Hesperolinon* species and included detailed comparisons of *Hesperolinon* to other Linoideae, but discounted the similarities of *Hesperolinon* to *Sclerolinon*, instead referring to subsection *Linopsis* ser. *Neo-Mexicana* as the closest relative of *Hesperolinon* based on features of the petal appendages and petal attachment to the staminal cup. Rogers (1975) and Xavier (1967) later cited morphological links between *Hesperolinon* and North American members of subsection *Linopsis* ser. *Linopsis*. We here find *Hesperolinon* and *Sclerolinon* to be most closely related to the *L. mexicanum* group, currently classified in subsection *Linopsis* ser. *Linopsis* (Fig 3.12).

**Character Evolution** — As indicators of evolutionary relationships and a basis for classification, Rogers attached particular importance to conditions of flower color, stylar fusion, heterostyly, the degree to which false septa penetrate the locules of the ovary, pollen morphology, leaf arrangement and morphology, and chromosome number (Rogers 1969, 1982). Derived features, such as the united styles and multiporate pollen that define subsection *Rigida*, appear to be good predictors of relationships in some cases, although some clades that appear in our results, like the lineage of South African *linums*, do not bear such easily recognizable potential synapomorphies. It appears that some of the derived states designated by Rogers (1982) have probably evolved multiple times from the ancestral conditions present in different lineages of subsection *Linopsis*. McDill et al. (2009) previously showed that heterostyly may have transitioned to homostyly multiple times in *Linum*. Evolutionary hypotheses for four additional characters — number of fruit segments, pollen morphology, development of false septa in the fruit, and stylar fusion — are presented in Fig. 3.13.

The number of segments into which the fruits of Linaceae dehisce is a product of the number of carpels and the number of sutures along which dehiscence occurs. The fruits of most *Linum* are capsules composed of five fused carpels, each carpel containing two seeds. Dehiscence of the capsule along both loculicidal and septicial sutures results in ten, one-seeded segments in most species of *Linum*. The number of fruit segments can change due to alteration in either the number of carpels or the number of sutures that dehisce. Both processes are considered together in Fig. 3.13 A, in which the number of capsule segments is mapped onto the phylogeny. Reduction in carpel number occurs in several lineages that have been recognized as segregate genera: in *Hesperolinon* (most species with three carpels, some with two (not sampled here)), *Sclerolinon* (two carpels, dehiscing along both suture types to produce four one-seeded nutlets), and *Radiola* (four carpels). The East African species *Linum keniense*, of section *Linopsis* subsect. *Keniense* (not sampled), also exhibits four carpels, and would likely represent an additional independent reduction in carpel number from the ancestral condition. In *Linopsis* subsect. *Rigida* ser. *Rigida*, the number of carpels is unchanged from the ancestral state, but the five carpels dehisce only along the loculicidal sutures, producing five capsule segments, each with two seeds (Rogers 1969). This change in dehiscence pattern (loss of dehiscence along the septicial sutures) coupled with retention of five carpels, results in the unique five-segmented fruit of series *Rigida*. *Linum sulcatum*, the single member of subsection *Rigida* ser. *Sulcata*, retains dehiscence along both suture types, resulting in ten-segmented capsules. Loss of septicial dehiscence also occurs in *Radiola* where, due to the co-occurrence of carpel reduction, the fruits dehisce into four segments.

Pollen grains of *Linum* are remarkably variable, exhibiting species-level variation in grain size and morphology, the size and shape of germinal pores, and in the size, shape, and morphology of exine processes (Xavier 1967, Rogers 1969, Xavier et al. 1980). The range of variation observed within *Linum* has been compared to that observed among genera in other families, and has been cited as an indication of “overly conservative” taxonomic treatment of *Linum* (Xavier 1967). Here we consider three broad categories of gross pollen morphology, referred to as tricolpate, colporate, and porate, based on the terminology of Xavier (1967). Tricolpate pollen is the most common type in section *Linopsis*, and was considered the “ancestral type” by Rogers (1982); the pollen grains are typically triangular in polar view, with three elongate germinal pores, or colpi (occasionally 4-6 colpi are present in 77tude77en77y species). Colporate pollen bear a larger number (~12) of elongate germinal pores, while porate pollen grains possess 20 or more circular pores. Pollen morphology exhibits five changes when mapped on our optimal phylogeny (Fig. 3.13 B), with “ancestral” tricolpate pollen transitioning to colporate in subsection *Linopsis* ser. *Virginiana* and to porate in subsection *Rigida*. In the series *Virginiana* clade, colporate pollen may have evolved in the common ancestor of the group, and been subsequently lost twice, or gained independently in three species.

The “false septa” often referred to by *Linum* taxonomists are extensions of the ovary wall penetrating to some extent into each locule, dividing the locule in half. The extent of these septa in the mature capsules varies on a continuous basis in section *Linopsis*, appearing to reach “completion” (with the septum forming a membrane that

completely divides the locule) in subsection *Rigida* ser. *Rigida* (all species), subsection *Linopsis* ser. *Virginiana* (in six of the seven species), subsection *Keniense* [not sampled here, but the condition was noted by Rogers (1982)], and in the segregate genera *Cliococca* and *Sclerolinon*. Complete septa may have evolved four times, with possible reversions to an incomplete state in some North American species of subsection *Linopsis* (Fig. 3.13 C).

In most species of Linaceae, the five styles are described as “free,” each distinct from the others from the point of attachment at the top of the ovary. In some species, the styles are fused together for a portion of their length, branching apart at some point well above the attachment to the ovary. Styler fusion is a hallmark of subsection *Rigida*, but also appears in several Mexican, South American, and Old World *linums* (Rogers 1982). Considering the occurrence of fused styles in *Linum* species worldwide, Rogers and Mildner (1976) hypothesized that this condition evolved multiple times in section *Linopsis*, and did not place too much importance on styler fusion in classifications or hypotheses of relationships. Rafinesque, on the other hand, used styler fusion as a basis for his segregate genus *Mesynium*, the name of which refers to the “fused middle.” He placed all of the fused-style yellow-flowered species he knew of in *Mesynium*, including an African species, *M. africanum* Raf. (= *Linum africanum* L.). When mapped on the molecular phylogeny, styler fusion exhibits as many as twelve independent gains, or four gains and seven reversions to free styles, in the most parsimonious reconstruction (Fig. 3.13 D). In light of our phylogenetic results, assignment of species to *Mesynium* based on styler fusion would clearly result in a polyphyletic genus.

While the derived character states designated by Rogers in many cases appear to have evolved multiple times from the ancestral conditions, his characterization of subsection and series *Rigida* as the most “advanced” lineage of section *Linopsis* is borne out by our results. Only in subsection *Rigida* are derived states of multiple characters present in most species, and at least one feature, reduction to only loculicidal dehiscence, appears to be a potential unique synapomorphy for series *Rigida*. While Rogers repeatedly noted the distinctiveness of subsection *Rigida*, and remarked that it might merit recognition as a genus, the existence of “intermediate” species such as *L. sulcatum* and occurrence of many of the “derived” states in other species of *Linum* led him to maintain this group as a subsection of *Linum* sect. *Linopsis* (Rogers 1969).

## CONCLUSION

The evolutionary scenario proposed by Rogers (1982) is shown to have been perhaps overly simplistic when examined in a phylogenetic framework, but his general view of character state “polarity” is essentially confirmed for a number of morphological features. However, Rogers’ classification scheme for *Linum* section *Linopsis*, based primarily on overall morphological similarity, is not supported by the molecular phylogenetic results. Several of Rogers’ subsections and series, those based at least in part on the presence of derived character states, such as subsection *Rigida* and subsection *Linopsis* ser. *Virginiana*, do correspond to supported clades in the molecular phylogenies. Other taxa, notably section *Linopsis*, subsection *Linopsis*, and series *Linopsis*, are

paraphyletic. We find nested among the lineages of *Linum* sect. *Linopsis* several taxa generally recognized at equal or higher rank (*Linum* sects. *Cathartolinum* and *Syllinum*, the segregate genera), further illustrating the need for a revised taxonomy for this lineage and for *Linum* in general.

The *Linum Rupestre*, *L. mexicanum*, and *L. schiedeana* species groups appear to be monophyletic, morphologically cohesive lineages that are moderately-to-well supported by our molecular data that have not been previously formally recognized in taxonomic treatments. The South American and South African species groups also appear to be distinct lineages, though morphological synapomorphies for these clades have not yet been discovered. While the phylogeny presented here does not completely resolve the evolutionary relationships among the species of *Linum* and its segregate genera, it is our hope that they will provide a basis for future rigorous comparative morphological study and taxonomic revision, and guide future phylogenetic sampling.

Table 3.1. Infraclassification of *Linum* sect. *Linopsis* (Rogers 1982). Species sampled for phylogenetic analyses in this study are indicated by asterisks (\*).

<i>Linum</i> sect. <i>Linopsis</i> (Rchb.) Engelm. in A. Gray		
Flowers four or five-merous, yellow (white to pink or purplish in a few spp.), petals separate, stigmas capitate or linear, sepals typically with marginal glands. Homostylous or heterostylous. Pollen tricolpate or multiporate. Approximately 94 spp.		
subsection 1: <i>Dichrolinum</i> (Planch.) Rogers		
Flowers five-merous, white to purplish, styles separate, stigmas capitate, Heterostylous or homostylous. Pollen tricolpate. $x=9$ . 2 spp.		
Mediterranean Region		
<i>L. suffruticosum</i> L. *	<i>L. tenuifolium</i> L. *	
subsection 2: <i>Halolinum</i> (Planch.) Rogers		
Flowers five-merous yellow, styles separate, stigmas linear, false septa partial, Heterostylous or homostylous. Pollen tricolpate. $x=10$ . 3 spp.		
Mediterranean Region		
<i>L. maritimum</i> L. *	<i>L. tenue</i> Desf. *	<i>L. trigynum</i> L. *
subsection 3: <i>Keniense</i> Rogers		
Flowers yellow, four-merous, styles separate, stigmas capitate, false septa complete. Homostylous. $x=?$ . 1 sp		
East Africa	<i>L. keniense</i> Fries	
subsection 4: <i>Linopsis</i> (Rchb.) Rogers		
Flowers five-merous, yellow, heterostylous or homostylous, styles separate or united, stigmas capitate, false septa partial to nearly complete, $x=9$ . Approximately 70 spp.		
series <i>Linopsis</i> (Rchb.) Planch.		
Homostylous (two spp. possibly secondarily heterostylous), styles separate or united, false septa partial, pollen tricolpate, $x=9$ . Approximately 59 spp.		
Mediterranean Region	Madagascar	<i>L. scoparium</i> Griseb. *
<i>L. strictum</i> L. *	<i>L. betsiliense</i> Baker	<i>L. smithii</i> Mildner
East Africa	<i>L. emirnense</i> Bojer	North America
<i>L. holstii</i> Engler ex Wilczek *	<i>L. marojejyense</i> (Humbert) Rogers	<i>L. arenicola</i> (Small) Winkler *
<i>L. volkensii</i> Engl. *	South America	<i>L. bahamensei</i> Northrop *
South Africa	<i>L. brevifolium</i> St. Hil. & Naud.	<i>L. cruciata</i> Planch. *
<i>L. acuticarpum</i> Rogers *	<i>L. burkartii</i> Mildner *	<i>L. flagellare</i> (Small) Winkler *
<i>L. adustum</i> E. Mey. Ex Planch. *	<i>L. carneum</i> Urb. *	<i>L. guatemalense</i> Benth. *
<i>L. aethiopicum</i> Thunb. *	<i>L. chamissonis</i> Schiede *	<i>L. gypsogenium</i> Nesom
<i>L. africanum</i> L. *	<i>L. cremnophyllum</i> Johnst. *	<i>L. lasiocarpum</i> Rose *
<i>L. brevistylum</i> Rogers	<i>L. erigeroides</i> St. Hil. *	<i>L. longipes</i> Rose *
<i>L. comptonii</i> Rogers *	<i>L. filiforme</i> Urb.	<i>L. mcvaughii</i> Rogers
<i>L. esterhuysenae</i> Rogers *	<i>L. littorale</i> St. Hil. *	<i>L. mexicanum</i> HBK *
<i>L. gracile</i> Planch. *	<i>L. macraei</i> Nenth. *	<i>L. modestum</i> Rogers
<i>L. heterostylum</i> Rogers *	<i>L. oligophyllum</i> Willd. Ex Schult. *	<i>L. nelsonii</i> Rose *
<i>L. pungens</i> Planch.	<i>L. organense</i> Gardn.	<i>L. orizabae</i> Planch. *
<i>L. quadrifolium</i> L. *	<i>L. palustre</i> Gardn.	<i>L. rupestre</i> (A. Gray) Engel. *
<i>L. thesioides</i> Bartl. *	<i>L. polygaloides</i> Planch.	<i>L. scabrellum</i> Planch. *

<i>L. thunbergii</i> E. & Z. *	<i>L. prostratum</i> Domb. Ex Lam. *	<i>L. schiedeanum</i> Schlecht. & Cham. *
<i>L. villosum</i> Rogers *	<i>L. ramosissimum</i> Gay	<i>L. tenellum</i> Schlecht. & Cham. *
<b>series <i>Mysorens</i> Rogers</b>		
Homostylous, styles united basally, false septa partial, pollen tricolpate, n=30. 1 spp.		
India	<i>L. mysorens</i> Heyne ex Benth.	
<b>series <i>Neo-Mexicana</i> (Small) Rogers</b>		
Homostylous, styles separate, false septa partial, pollen tricolpate, x=13. 3 spp.		
Western North America		
<i>L. kingii</i> S. Watson*	<i>L. neomexicanum</i> HBK*	<i>L. pringlei</i> S. Watson*
<b>series <i>Virginiana</i> (Small) Rogers</b>		
Homostylous, styles separate, false septa partial to complete, pollen tricolpate or few-porate, x=18. 7 spp.		
Eastern U.S.		
<i>L. floridanum</i> (Planch.) Trel. *	<i>L. medium</i> (Planch.) Britton *	<i>L. westii</i> Rogers *
<i>L. intercursum</i> Bicknell *	<i>L. striatum</i> Walter *	
<i>L. macrocarpum</i> Rogers	<i>L. virginianum</i> L. *	
<b>subsection 5: <i>Rigida</i> (Small) Rogers</b>		
Flowers yellow, five-merous, styles variously united, stigmas capitate, false septa partial to complete, pollen multiporate, x=15. 15 spp.		
<b>series <i>Rigida</i> (Small) Rogers</b>		
Styles united to above the middle, false septa complete. 15 spp.		
Central and southwestern North America		
<i>L. alatum</i> (Small) Winkler *	<i>L. compactum</i> A. Nelson *	<i>L. puberulum</i> (Engel.) Heller *
<i>L. aristatum</i> Engel. in Wisl. *	<i>L. elongatum</i> (Small)Winkler *	<i>L. rigidum</i> Pursh *
<i>L. australe</i> Heller *	<i>L. hudsonioides</i> Planch. *	<i>L. subteres</i> (Trel.) Winkler *
<i>L. berlandieri</i> Hooker *	<i>L. imbricatum</i> (Raf.) Shinners *	<i>L. vernale</i> Wooton *
<i>L. carteri</i> Small *	<i>L. lundellii</i> Rogers *	<i>Linum</i> sp. nov.
<b>series <i>Sulcata</i> (Small) Rogers</b>		
Styles united to below the middle, false septa partial. 1 spp.		
North America	<i>L. sulcatum</i> Riddel *	



Table 3.2. Primers used to amplify and sequence the *matK* gene from yellow-flowered *Linum* and outgroups.

Primer Name	Sequence (5'-3')	Taxonomic Utility
<i>matK</i> 1F	ACT GTA TCG CAC TAT GTA TCA	all
<i>matK</i> 1320R	GAT CCG CTA TAA TAA TGA GA	all
<i>matK</i> 595F	TGG ATA AAA GAT CCC GCT TG	Yellow-flowered taxa
<i>matK</i> 595R	CAA GCG GGA TCT TTT ATC CA	Yellow-flowered taxa
<i>matK</i> 819F	TCA GTT TCT TCT TGA GCG AAA G	Yellow-flowered taxa
<i>matK</i> 819R	CTT TCG CTC AAG AAG AAA CTG A	Yellow-flowered taxa
<i>matK</i> 819Fb	TAG CTT YYT TCT TGA GCG AAA T	Blue-flowered taxa/OG
<i>matK</i> 819Rb	ATT TCG CTC AAG AAR RAA GCT A	Blue-flowered taxa/OG

Table 3.3: Data matrix characteristics. NTax = number of terminal taxa, Nchar = total number of base pairs in aligned matrix. Number included = number of base pairs included in analyses (after exclusion of questionably-aligned sites). G1 = G1 statistic, determined from evaluation of one hundred thousand random trees in PAUP. Constant = # of constant characters. P.i. = number of parsimony informative characters. Model=substitution model type selected.

Matrix:	<i>matK</i>	<i>trnK</i> intron	<i>ndhF</i>	<i>trnL-F</i>	CPL Combined	ITS	CPL + ITS
NTax	135	105	132	129	136	135	122
NChar	1369	409	490	1309	3577	777	4354
Number included	1154	405	433	753	2745	624	3369
G1	-1.53	-0.69	-1.59	-1.34	-1.40	-0.51	-0.57
constant	721	257	317	615	1910	310	2509
p.i.	312	75	94	215	696	241	635
Model	GTR+ $\Gamma$	F81+ $\Gamma$	GTR+ $\Gamma$	GTR+ $\Gamma$	GTR+ $\Gamma$	GTR+ $\Gamma$	GTR+ $\Gamma$

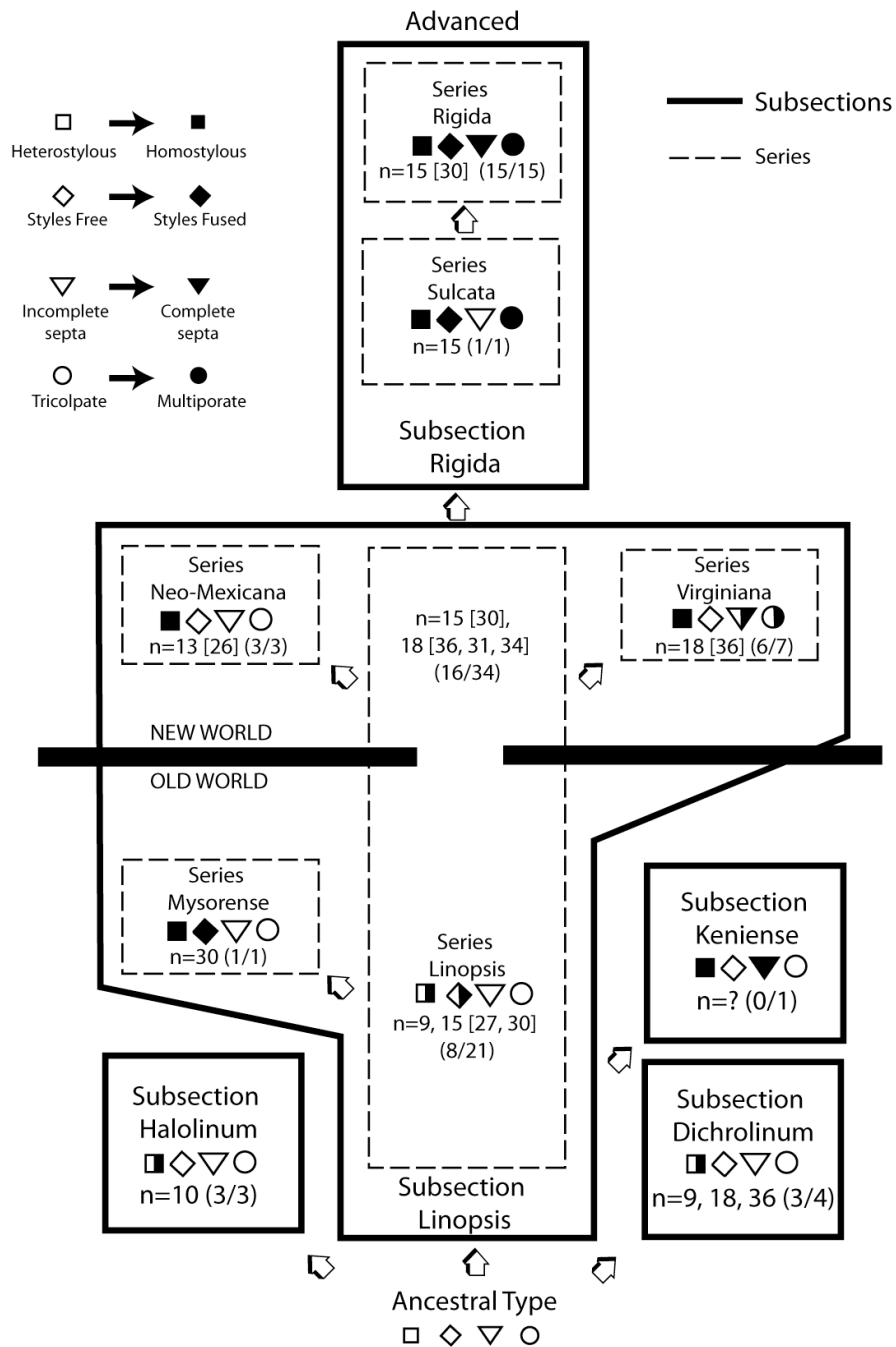
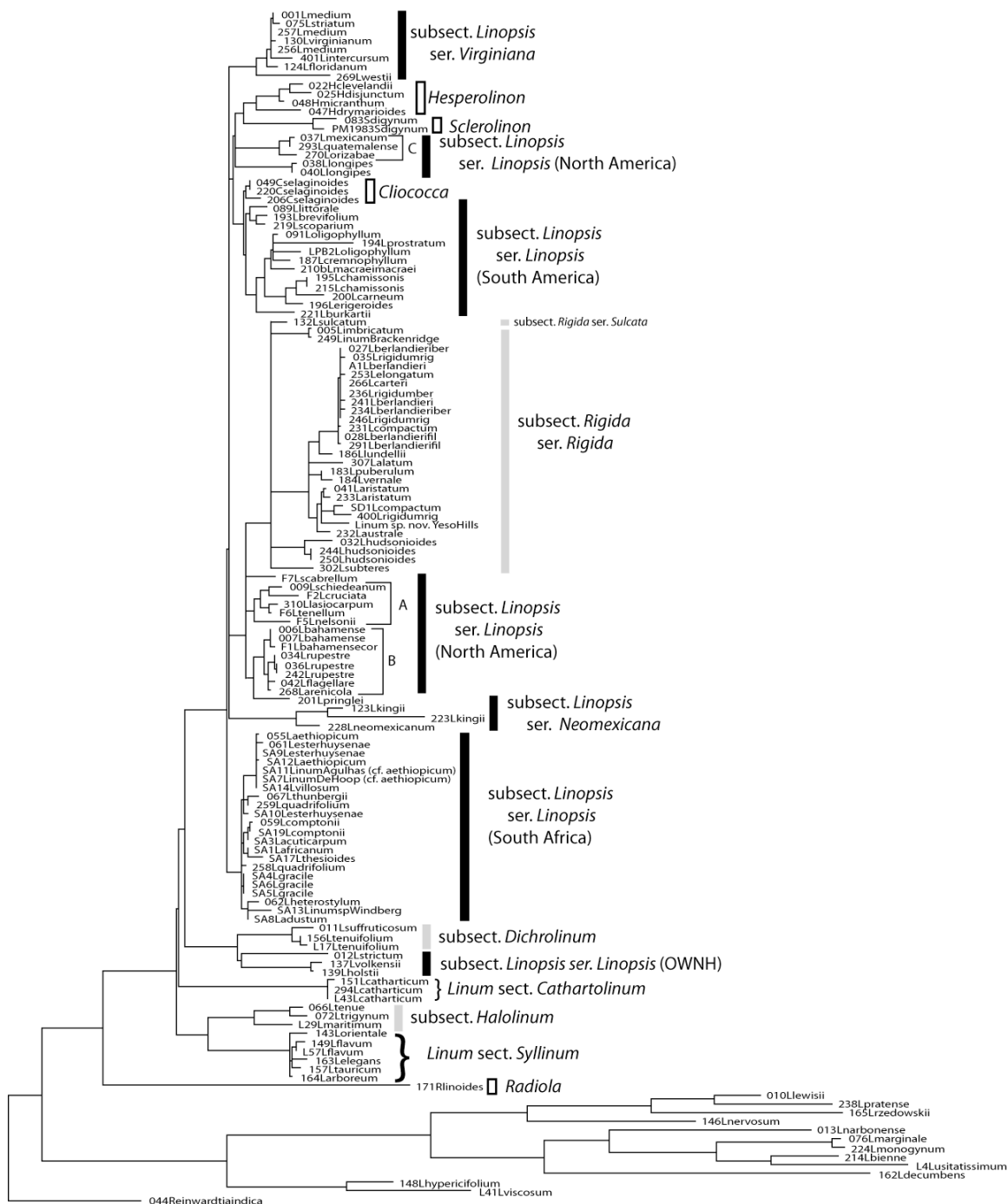


Fig. 3.1. Hypothesized evolutionary relationships among the various subgroups of *Linum* sect. *Linopsis* adapted from Rogers (1982, Fig. 1). Open shapes indicate “primitive” character states designated by Rogers, filled shapes the “derived” state.



- 0.001 substitutions/site

Fig. 3.2. Maximum likelihood topology for CPL matrix (-lnL=15848.70). Braces = sects. Cathartolinum and Syllinum. Shaded bars indicate the subsectional classifications (Table 3.1): Grey = sect. *Linopsis* subsections. Dichrolinum, Halolinum, and Rigida. Black = subsection *Linopsis*. Open = segregate genera. Lettered clades: A = *Linum schiedeanum* group, B = *Linum rupestre* group, C = *Linum mexicanum* group.



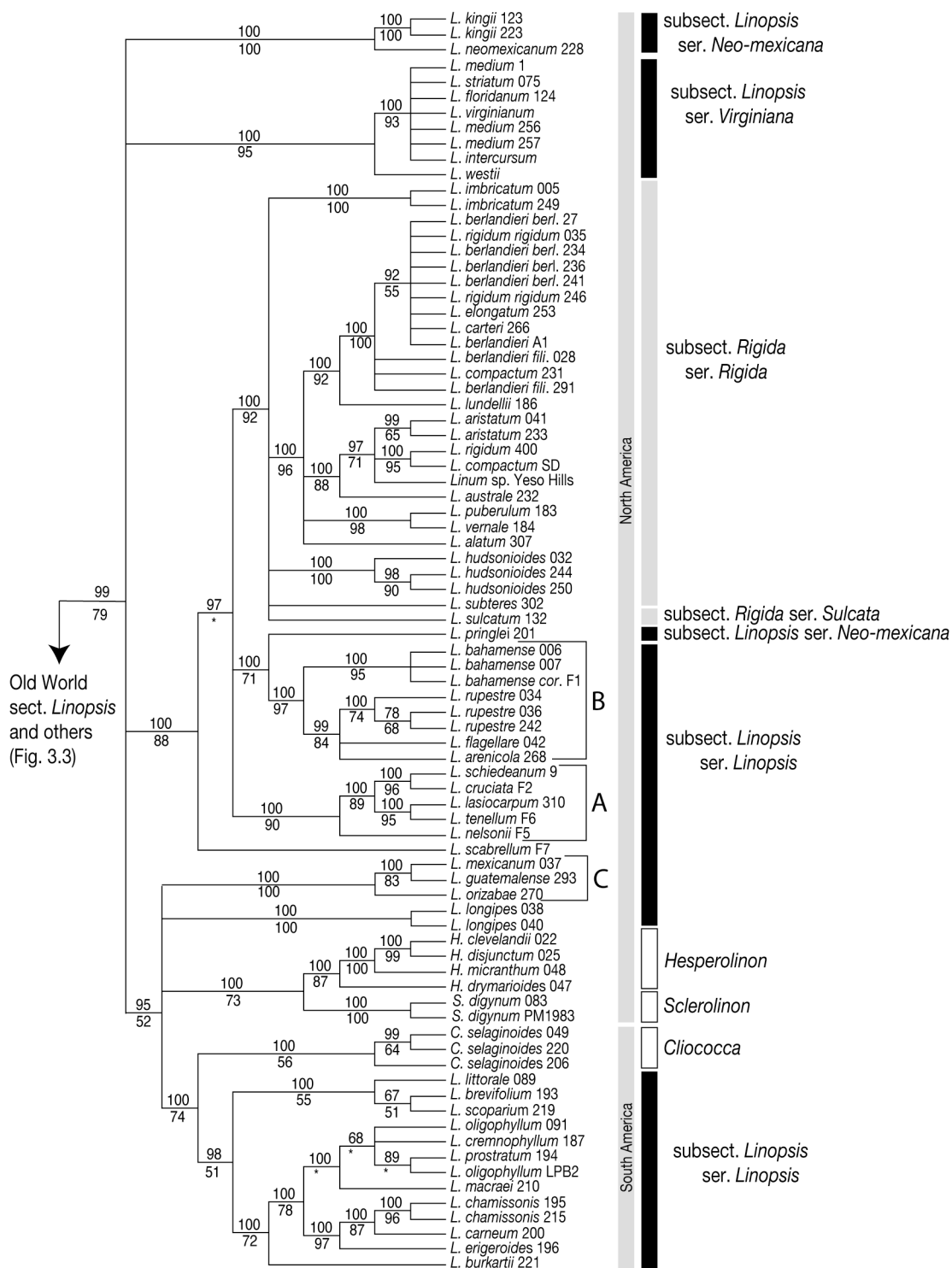


Fig. 3.4. Continuation of Fig 3.3, Part 2: New World taxa. The informal species groups referred to in the text are indicated by lettered brackets: A: *Linum schiedeanum* group, B: *Linum rupestre* group, C: *Linum mexicanum* group.

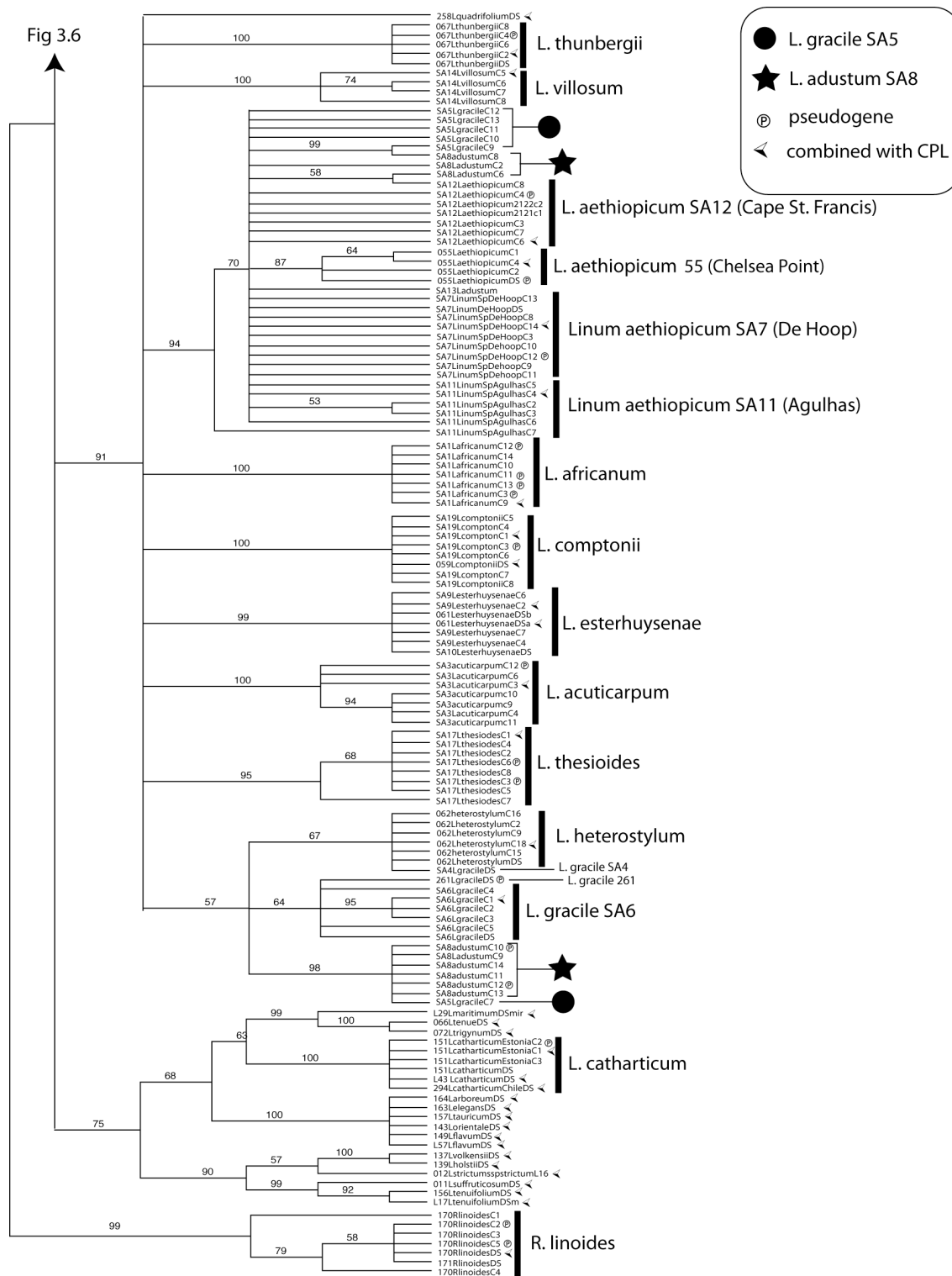


Fig. 3.5. ITS Clone sequence ML bootstrap 50% majority-rule consensus, Part 1. Nodes with less than 50% support are collapsed.

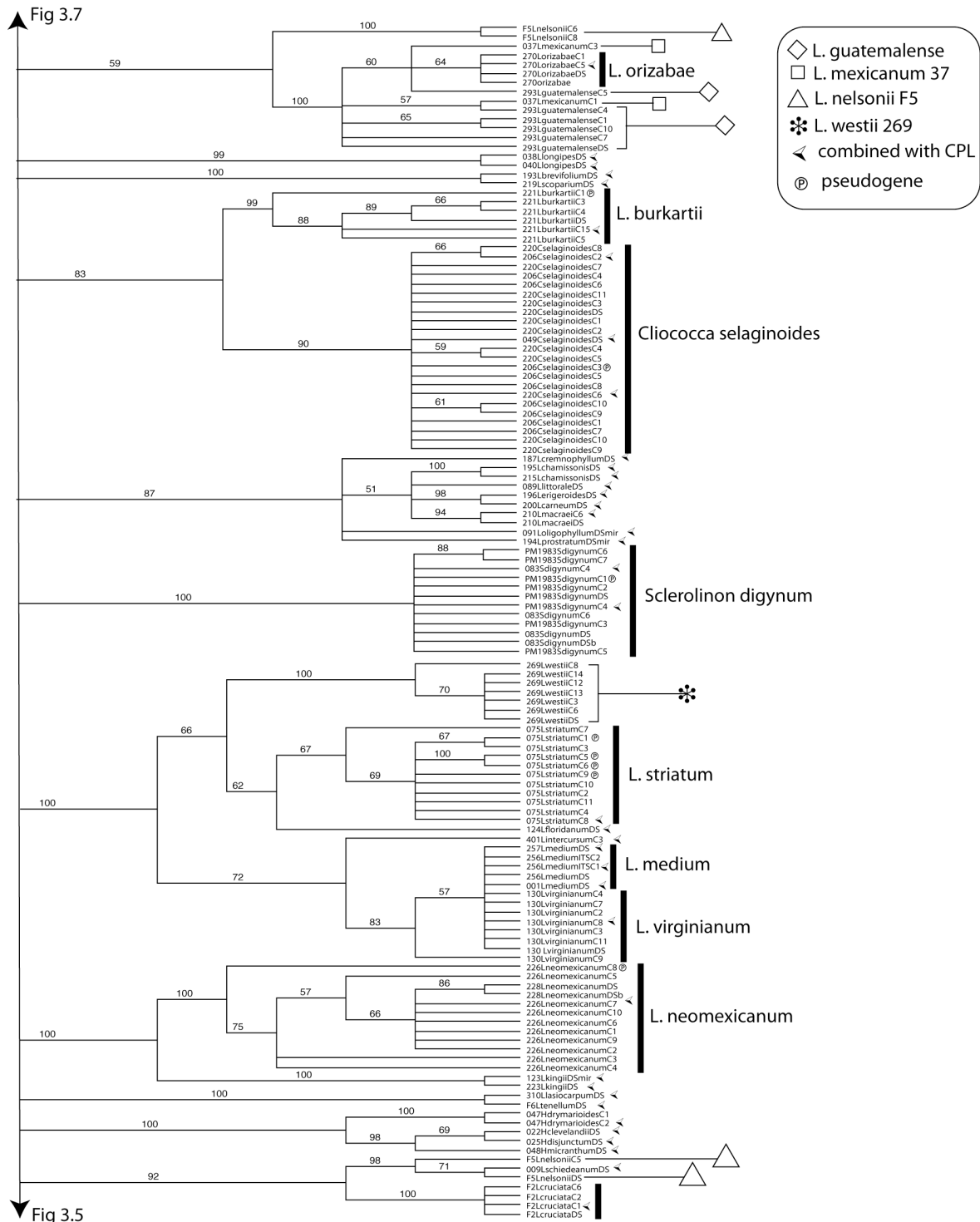


Fig. 3.6. ITS Clone sequence ML bootstrap 50% majority-rule consensus, Part 2. Nodes with less than 50% support are collapsed.



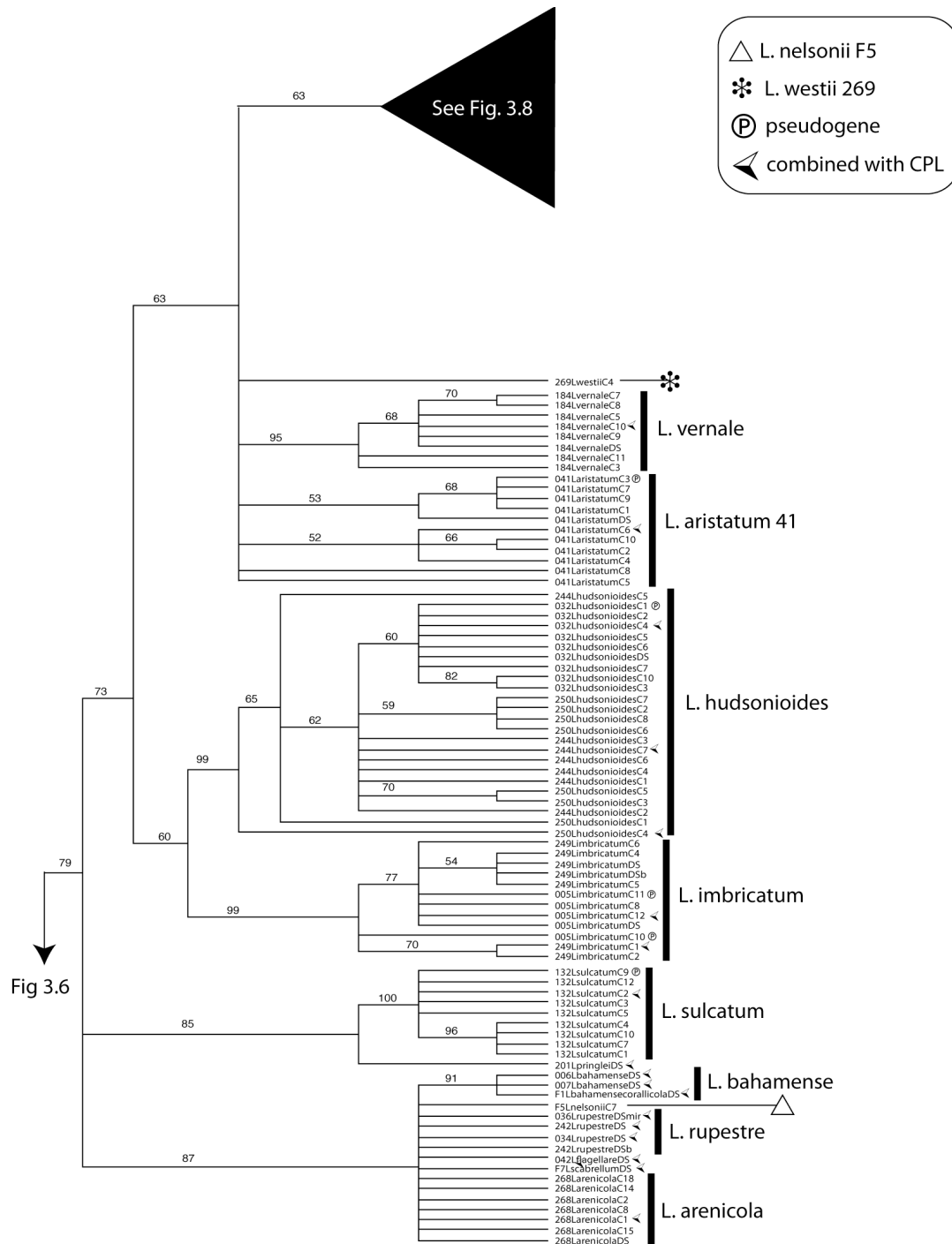


Fig. 3.7. ITS Clone sequence ML bootstrap 50% majority-rule consensus, Part 3. Nodes with less than 50% support are collapsed.

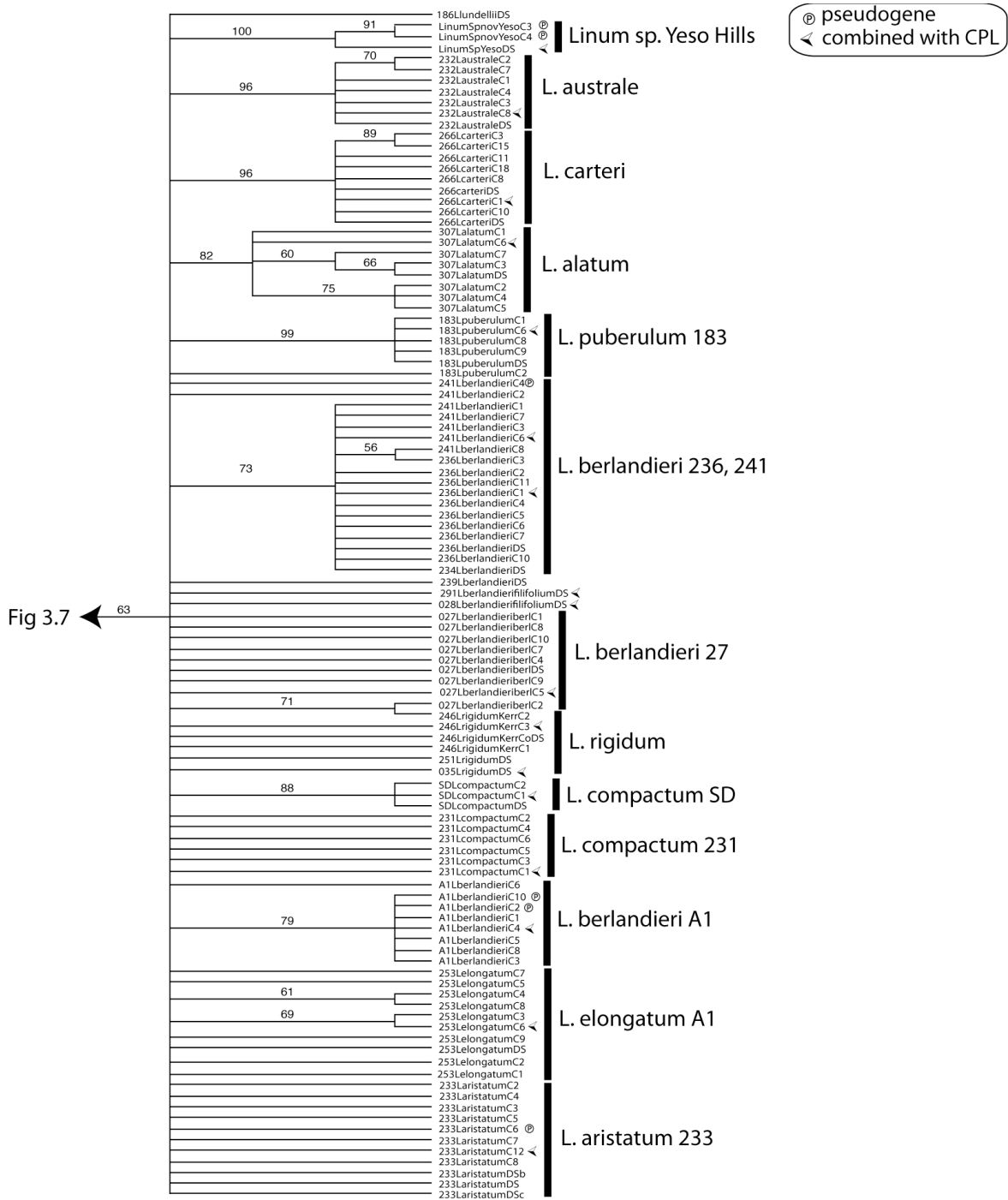
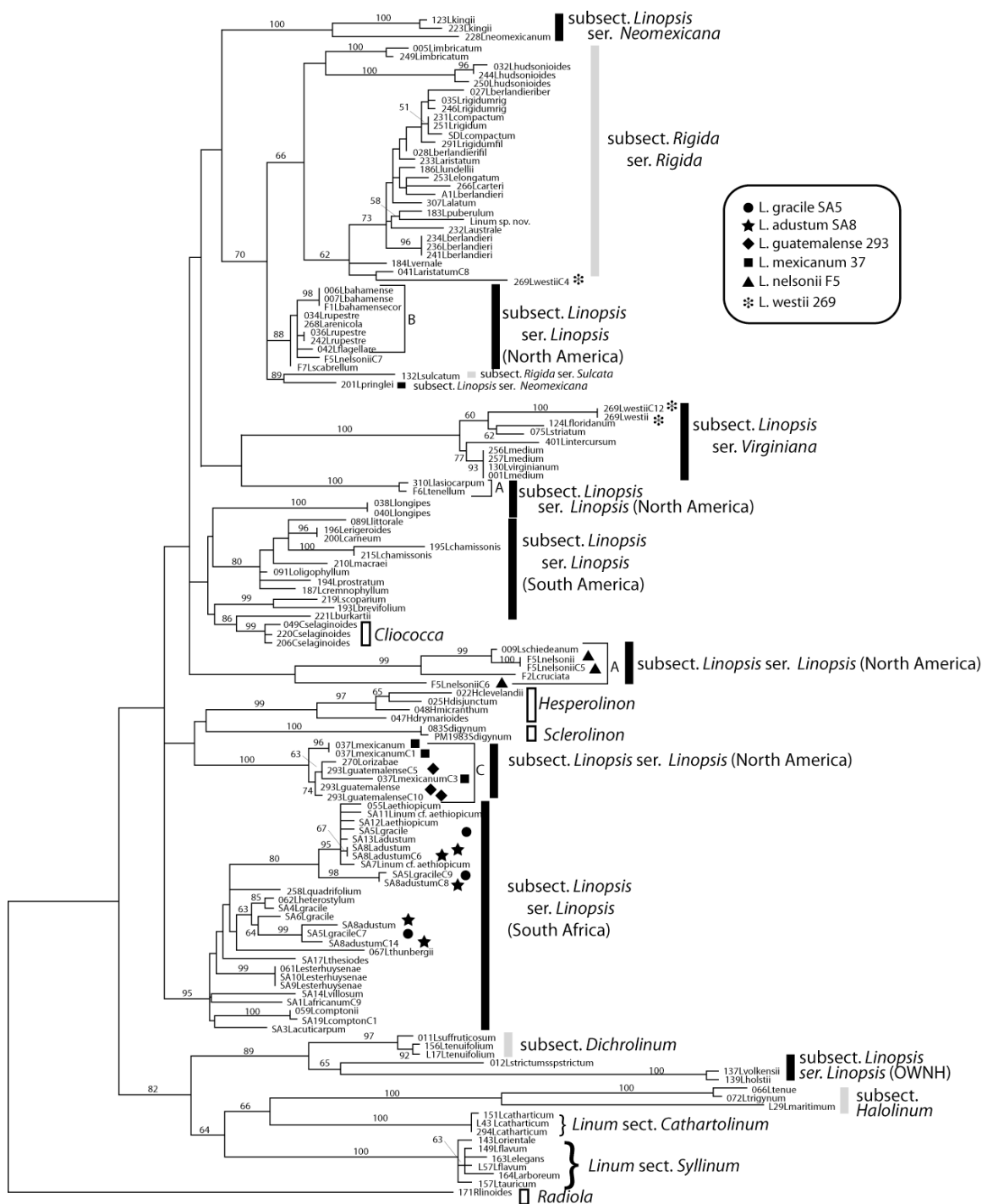


Fig. 3.8. ITS Clone sequence ML bootstrap 50% majority-rule consensus, Part 4. Nodes with less than 50% support are collapsed.



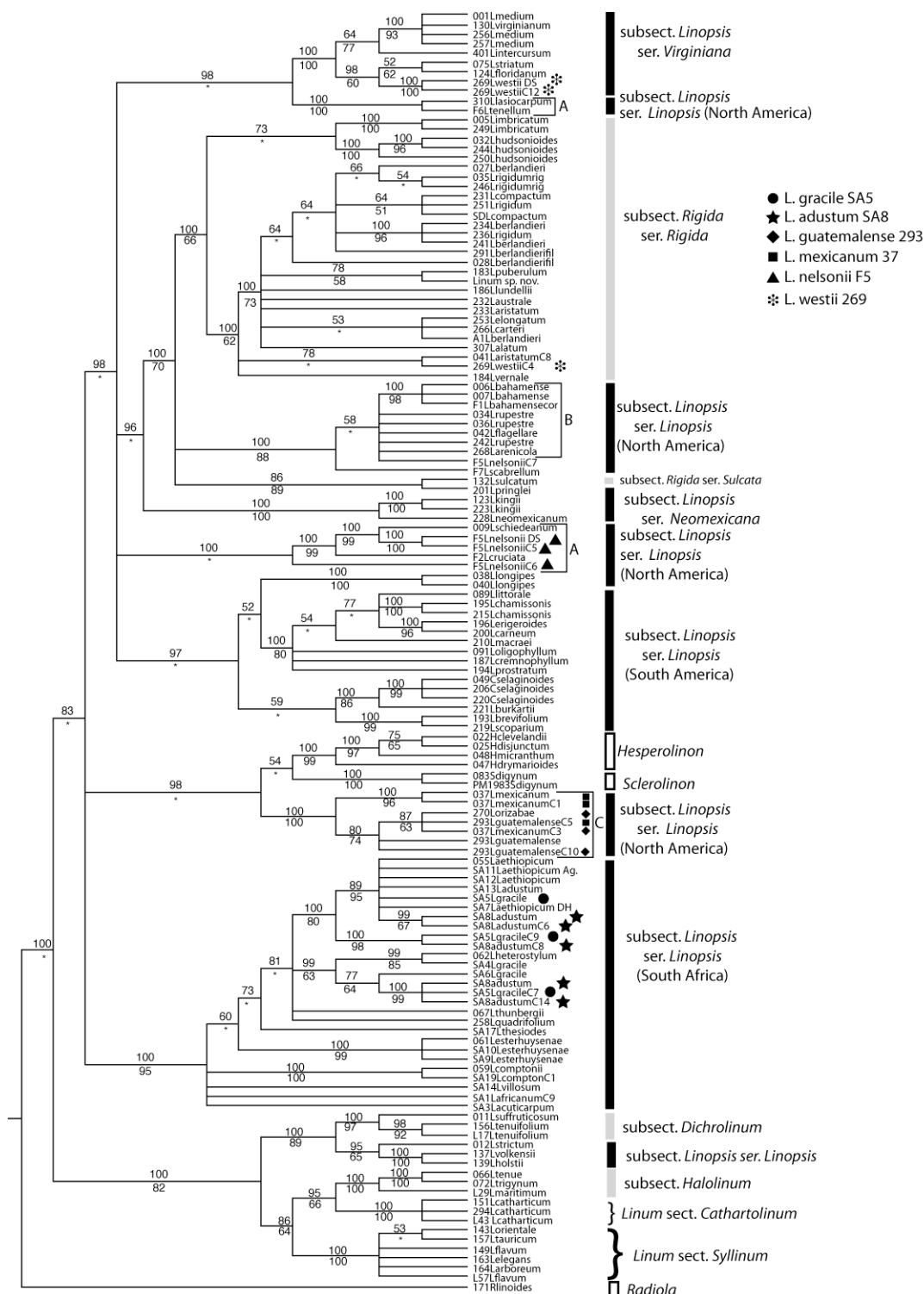


Fig. 3.10. Majority-rule consensus from Bayesian analysis of ITS. Posterior probabilities (as %) above branches, ML bootstrap below (<50% indicated by \*). A=*L. schiedeanum* group, B=*L. rupestre* group, C=*L. mexicanum* group.

0.005 substitutions/site

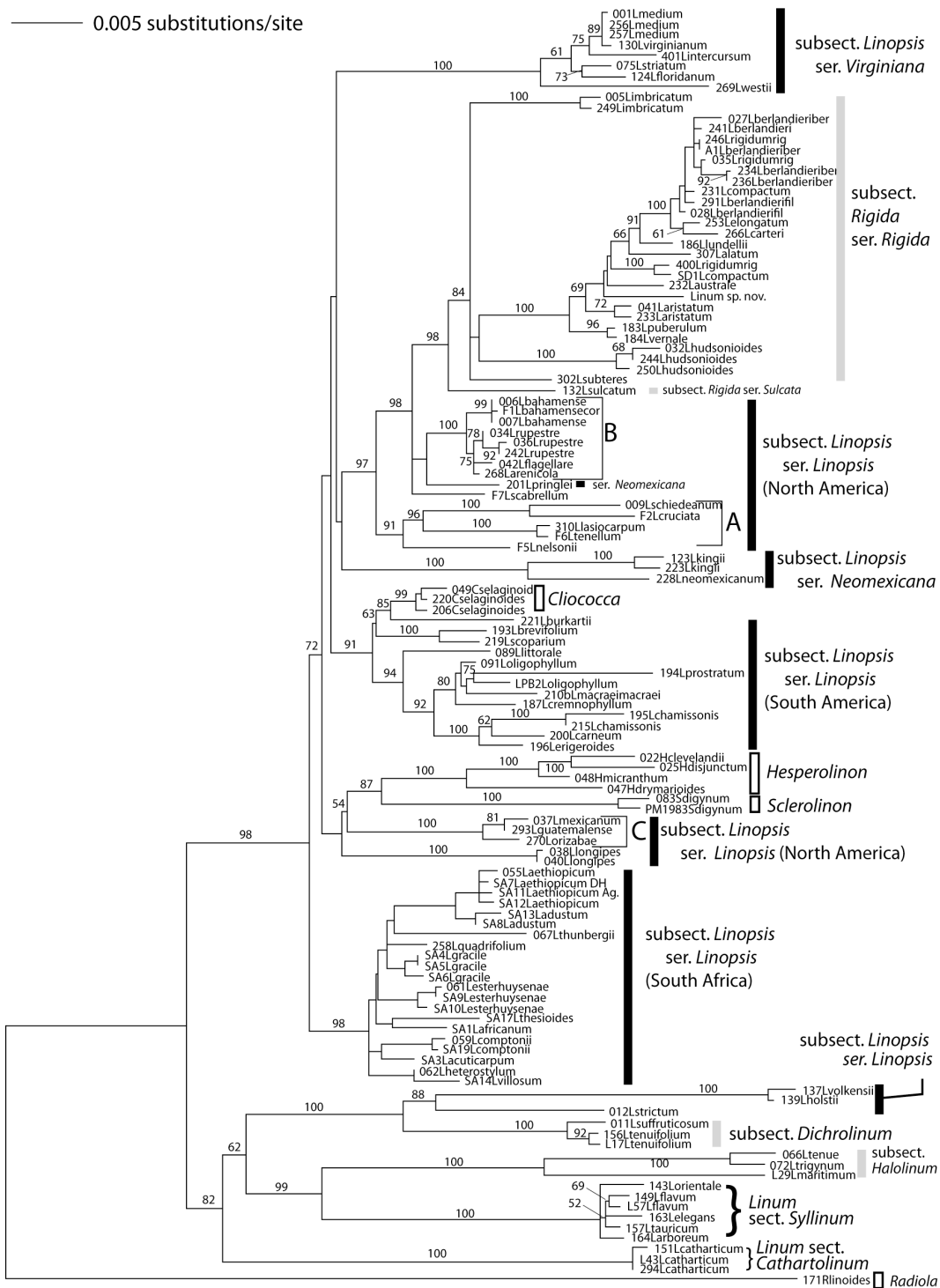


Fig. 3.11. Maximum-likelihood topology ( $-\ln L = 18245.7291$ ) for combined CPL+ITS matrix, with ML bootstrap percentages above branches ( $<50\%$  not shown). Clade A = *Linum schiedeanum* group, B = *Linum rupestre* group, C = *Linum mexicanum* group.

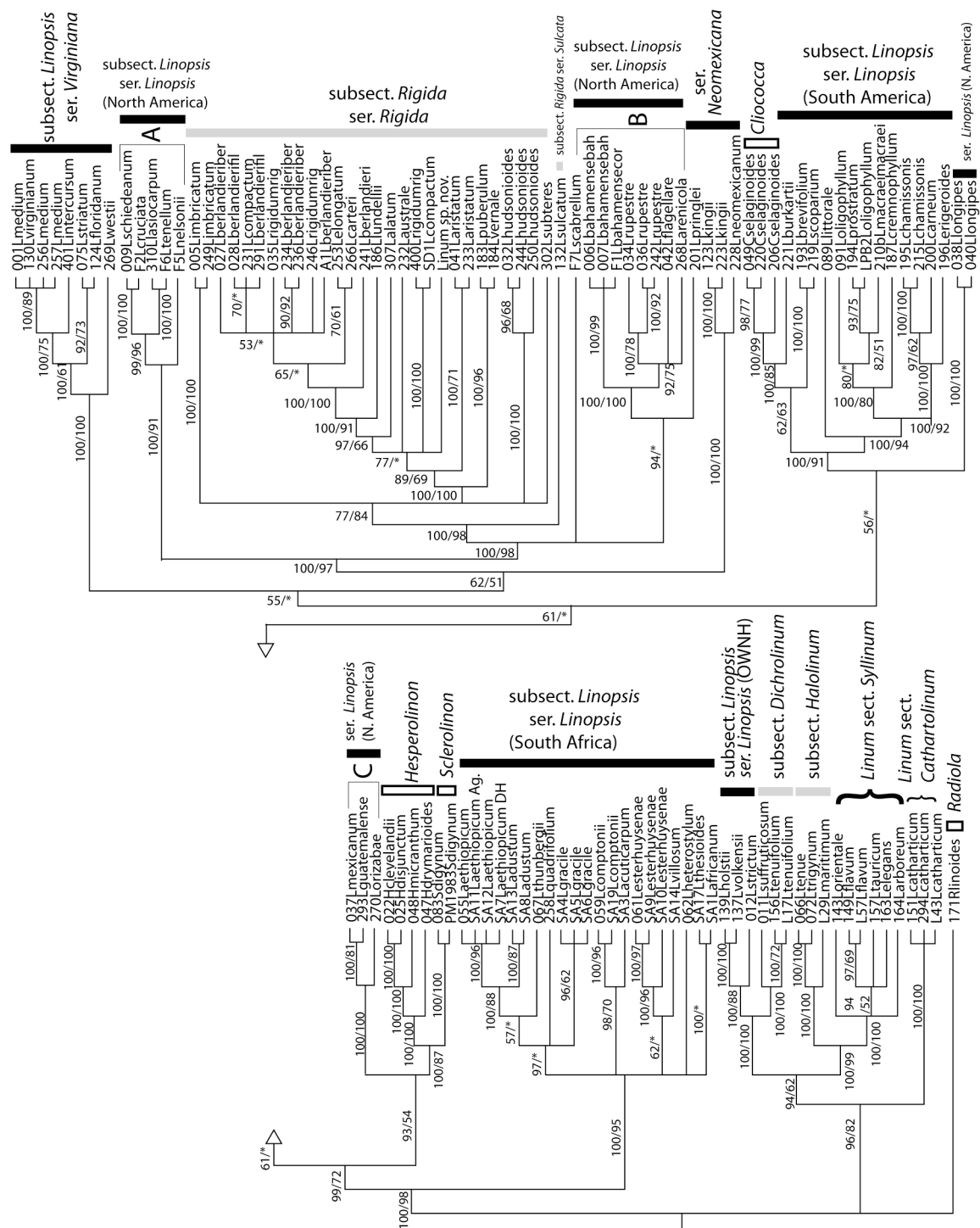
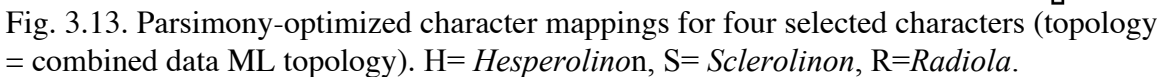


Fig. 3.12. Majority-rule consensus from Bayesian analysis of combined CPL+ITS data; posterior probabilities (as percentages)/ML bootstrap percentages (values <50% indicated by \*). A = *Linum schiedeanum* group, B = *Linum rupestre* group, C = *Linum mexicanum* group.



## Chapter 4: On the Origins of Multiple Intercontinental Disjunctions in Yellow-Flowered *Linum* (Linaceae)

### ABSTRACT

Yellow-flowered lineage of *Linum* (including the segregate genera *Cliococca*, *Hesperolinon*, *Radiola*, and *Sclerolinon*) has diversified on four continents: Eurasia, Africa, North America, and South America. Potential historical biogeographic scenarios for the establishment of species groups on each continent are evaluated in the context of molecular phylogenetic hypotheses based on nuclear (ITS) and chloroplast (*matK*, *ndhF*, *trnK* 3' intron, and *trnL-F* intron and spacer) data and sampling from the majority of species from each region. All New World species form an apparently monophyletic group, suggesting a single Amphi-Atlantic disjunction, although statistical tests can not rule out the possibility of two trans-Atlantic events. The timing of this Amphi-Atlantic disjunction, based on relaxed Bayesian molecular clock analysis, indicates migration across the fragmenting North Atlantic land bridge in the Miocene may potentially explain this disjunction. Both the South American and South African species groups are well supported as monophyletic, indicating single colonization's of both southern continents from Northern Hemisphere ancestry. Establishment of these amphitropical disjunctions are also estimated to have occurred in the Miocene. We find no compelling evidence for recent intercontinental dispersal in this lineage of *Linum*, in accordance with the probably low dispersal potential exhibited by their dehiscent capsular fruits and large, smooth seeds.

### INTRODUCTION

Best known as the genus of the cultivated flax, *L. usitatissimum* L., *Linum* includes nearly 200 species of annuals and suffrutescent perennials distributed throughout



temperate and tropical regions of six continents. Considered as a whole, in its current circumscription, *Linum* has a subcosmopolitan distribution (Thorne 1972), but the two major lineages of *Linum* exhibit drastically different distributional patterns. The blue-flowered flaxes of *Linum* sects. *Linum* and *Dasylinum* (shown to comprise a monophyletic lineage in McDill et al. 2009) are essentially confined to Eurasia, with only a few species of section *Linum* native to other continents: three species in North America (Rogers 1984), and two in Australia and New Zealand (Rogers and Xavier 1971). In remarkable contrast, diverse species groups of yellow-flowered flaxes (*Linum* sect. *Linopsis*) exist in Eurasia, North America, South America, and Africa, with additional isolated species in the Galapagos islands, Madagascar, and India, but none native to Australia or New Zealand (Rogers 1982). Recent and ongoing molecular phylogenetic analyses have shown the species of section *Linopsis* to be part of a lineage that includes *Linum* sections *Cathartolinum* and *Syllinum*, and the segregate genera *Cliococca*, *Hesperolinon*, and *Sclerolinon* (McDill et al. 2009). This lineage has been loosely referred to as the “yellow-flowered clade” of *Linum*, and is consistently found to be sister to the monotypic genus *Radiola* Roth. (McDill et al. 2009, and Chapters 2 and 3 of this dissertation). The geographic distribution of the group is depicted in Fig. 4.1, with the approximate number of species indicated for each area based on floristic and taxonomic treatments. This distribution implies that multiple trans-Atlantic and trans-tropical disjunctions have taken place in the course of the diversification of yellow-flowered *Linum*.

Botanists have been intrigued by such intercontinental disjunctions for over 150 years (Gray 1846, Hooker 1860, Thorn 1972, Raven and Axelrod 1974), and several general biogeographic hypotheses have been proposed to explain amphi-Atlantic and amphitropical disjunctions in plants. In the Southern Hemisphere, transatlantic disjunctions between Africa and South America are sometimes considered products of vicariance due to the breakup of Gondwana (Raven and Axelrod 1974). In the Northern Hemisphere, transatlantic disjunctions between North American and Eurasia might result from Gondwanan vicariance followed by dispersal or migration into the Northern Hemisphere (Davis et al. 2002a), or migration across ancient North Atlantic or Beringian land-bridge connections followed by isolation when the bridges became discontinuous (Thorne 1972, Raven and Axelrod 1974, Wen and Ickert-Bond 2009). This Boreotropic Hypothesis (Wolfe 1975, Tiffney 1985) postulates the movement of tropical and warm-temperate taxa across a North Atlantic land bridge during the Eocene and Oligocene (55 to 25 mya). This idea has been used to explain modern amphi-Atlantic disjunctions of taxa whose divergence times or phylogenetic relationships are found to be incompatible with Gondwanan vicariance (Lavin and Luckow 1993, Davis et al. 2004, Weeks et al. 2005, Milne 2006). Long-distance dispersal — mediated by animals, wind, or floating rafts of debris — provides another potential mechanism for the establishment of amphi-Atlantic disjunct lineages in either hemisphere (Raven 1963, Thorne 1972, Houle 1998, Givnish and Renner 2004, Renner 2004).

Amphi-tropical disjunctions, in which closely-related species are found in temperate areas of the Northern and Southern hemispheres but are absent from the

intervening tropical regions, are observed in numerous genera in both the Old and New World, with New World amphitropical disjuncts receiving the most attention from botanists (Thorne 1972, Wen and Ickert-Bond 2009). Most cases, particularly those involving disjunct species or species pairs, are generally considered the products of relatively recent ( $<2$  mya) long-distance seed dispersal, with migrating shorebirds frequently implicated as potential agents of transportation (Raven 1963, Wen and Ickert-Bond 2009). More ancient dispersals ( $>15$  mya) have also been invoked for some taxa, such as *Ephedra* (Ickert-Bond et al. 2009), where divergence times between North and South America greatly predate the closure of the Panamanian isthmus and the establishment of migratory pathways. Amphitropical distributions could also be the product of a historically continuous distributions of species which became disjunct due to extinction in the tropics (Raven 1963), though this scenario has not been supported by findings from molecular phylogenetic studies of amphitropical disjuncts to date (Wen and Ickert-Bond 2009).

Few specific references have been made to potential biogeographic histories for Linaceae or *Linum*. Based on Paleocene occurrences of *Ctenolophon*-type pollen fossils in Africa and South America (Germeraad et al. 1968), Raven and Axelrod (1974) concluded that the amphi-Atlantic distribution of both subfamilies of Linaceae could be a result of vicariance due to the breakup of Gondwana. Though *Ctenolophon* has been considered a member of Linaceae in the past (Winkler 1931), molecular phylogenetic studies (Savolainen et al. 2000, Wurdack and Davis 2009) have shown that placement was probably incorrect, so its occurrence can not be used as direct evidence for the age of

Linaceae, and does not establish the presence of Linaceae in South America and Africa prior to the breakup of Gondwana. The earliest known fossils of Linaceae are *Linum* pollen grains from northeastern Spain, dated to the late Eocene, between 33.9 and 37.2 million years ago (Cavaghetto and Anadón 1996). As discussed in Chapter 2, the fossil pollen appears most similar to either the *Linum catharticum* or *Linum austriacum* pollen types defined by Punt and Den Breejen (1981). The fossil record thus establishes the presence of *Linum* in Europe in the late Eocene, but fossil occurrences from other continents and other genera of Linaceae are unknown.

In the course of revisionary work on *Linum* sect. *Linopsis*, both Rogers and Mildner noted possible relationships between Eurasian, African, and New World linums based on morphological similarities. Rogers proposed that *Linum* section *Linopsis* (and *Linum* generally) originated and initially diversified in Eurasia, and that the African and American species groups were ultimately derived from Eurasian ancestry (Rogers 1982). This was essentially confirmed by McDill et al. (2009), who found that lineages of Old World Northern Hemisphere Linoideae form a paraphyletic grade that subtends the African and New World lineages. Rogers also cited the occurrence of a few species of section *Linopsis* in Tropical East Africa (e.g., *Linum keniense*, *L. volkensii*, and *L. holstii*) as evidence of a biogeographic connection between South Africa and the Mediterranean region (Rogers 1982), an idea tentatively refuted by the results of McDill et al. (2009), which showed the four South African taxa sampled having an origin independent from the East African taxa (represented by *L. volkensii*).

Rogers cited morphological similarities of Mexican species of *Linum* to South African *Linum quadrifolium* Thunb. as possible evidence for a South African origin of North American *Linum* (Rogers 1969). While revising the taxonomy of South American *Linum*, Mildner and Rogers further discussed morphological similarities of “primitive” South American and Mexican flaxes to African taxa, suggesting close relationships among species on all three continents and the possible derivation of all New World members of section *Linopsis* from South African ancestry (Mildner 1971, Mildner and Rogers 1978). Although potential African ancestry of New World linums has been suggested repeatedly, no specific time frame or mechanism for the putative intercontinental disjunctions were proposed. The morphological similarity of species on the different continents could alternatively be explained as convergences or retained plesiomorphic states rather than indicators of recent common ancestry.

We can envision several historical scenarios by which yellow-flowered *Linum* could have attained its current distribution, each scenario with certain phylogenetic consequences. Three simplified scenarios and hypothetical phylogenies exhibiting the concomitant “expected” sets of relationships are presented in Fig. 4.2. A scenario fitting Rogers’ morphology-based hypothesis is shown in Fig. 4.2 A, with North American and South American taxa comprising a monophyletic group that is nested within (derived from) an African lineage. If New World flaxes originated from South African ancestry, we could expect the South African group to be non-monophyletic. Figure 4.2 B represents a scenario involving a northern hemisphere trans-Atlantic disjunction, which would be consistent with the movement of *Linum* across a North Atlantic land bridge or

dispersal from Europe to North America. In this scenario, amphitropical disjunctions are the product of independent colonizations of South America and South Africa from North America and Eurasia, respectively. Testable expectations of this scenario include North American non-monophyly (due to the derivation of South American *Linum* from within the North American lineage), New World monophyly, and South American monophyly. Figure 4.2 C represents an example of a more complex scenario, with independent Northern and Southern trans-Atlantic events, and independent Old and New world amphitropical events. In this scenario, the testable expectations would include the non-monophyly of all of the continental species groups.

In addition to topological expectations, these geology-linked biogeographic scenarios impose temporal expectations on the establishment of disjunct lineages. As mentioned above, Gondwanan vicariance requires the occurrence of the common ancestor of the disjunct lineages to be present prior to the separation of South America and Africa, which was well established by the late Cretaceous, approximately 100 mya (Sanmartin and Ronquist 2004). The North Atlantic land bridge is estimated to have provided a continuous migration pathway for temperate or warm-adapted plants from the early to mid Eocene (approximately 55 mya to 45 mya; Tiffney 1985, Tiffney and Manchester 2001). The land bridge became discontinuous during the latter half of the Eocene, and temperature, moisture, and light availability also becoming limiting factors, although the potential for migration by island hopping may have been possible as recently as 15 mya (Tiffney and Manchester 2001, Milne 2006, Bell 2007). The Bering

land bridge was also an available pathway for migration from Eurasia to North America from at least the early Eocene until the late Miocene (as recently as 5 mya).

More recent long-distance dispersals might also provide an explanation for the distribution of extant *Linum* species, but the genus presents challenges for dispersal scenarios as well. The fruits of all flaxes are dehiscent capsules and the seeds are large (ranging from 1 to 5 mm in length, depending on the species) and smooth. Other than the seeds of some species exuding a gelatinous matrix (and thus perhaps becoming sticky) when wet (pers. obs.), neither the fruits or seeds of *Linum* have any obvious adaptations for dispersal, so their potential for long-distance dispersal is also in question.

Our goal here is to utilize phylogenetic hypotheses based on chloroplast and nuclear DNA sequence data from an extensive sample of *Linum* species (Chapter 3) to investigate history of intercontinental disjunctions in the yellow-flowered lineage of *Linum*. To this end, we employ several different methods to compare the phylogenetic results with the expectations of different biogeographic scenarios. Parsimony-based optimizations of continental distributions on molecular phylogenies are used to identify the minimal number and direction of intercontinental disjunctions that may have taken place in this lineage. Divergence time estimates for the yellow-flowered clade are used to assess compatibility of the establishment and diversification of disjunct lineages with geological factors such as land bridge continuity. Finally we use a statistical method referred to as the SOWH test (Goldman et al. 2000) to discriminate among different potential biogeographic scenarios based on their predictions regarding the monophyly of different lineages of yellow-flowered linums.

## MATERIALS AND METHODS

**Taxon Sampling** — In the previous phylogenetic study of yellow-flowered *Linum* (Chapter 3), specimens were selected to represent the taxonomic diversity of this lineage on the four continents where the majority of species occur. While the attempt was made to accomplish complete taxon sampling (based on the most recent taxonomic and floristic treatments) for North American, South American, and African linums, viable DNA extracts for certain species were unavailable due to the unavailability of sample-able specimens or apparent degradation of DNA in specimens of advanced age. Other taxa (e.g. additional members of *Linum* sect. *Syllinum*) were not sampled because of their membership in geographically-restricted lineages we believe to be monophyletic based on earlier studies. A complete account of missing taxa, noting their geographic distributions, follows.

From Eurasia, the unsampled taxa of *Linum* section *Linopsis* are *L. corymbulosum*, *L. setaceum* (both Mediterranean species), and *L. mysorens* (endemic to India and Sri Lanka). Only five of an estimated 37 species of *Linum* section *Syllinum* were sampled; this section was found to be monophyletic as sampled by Vasilev et al. (2008) and McDill et al. (2009). Additional sampling of section *Syllinum* is unlikely to alter our understanding of the historical biogeography of the yellow-flowered lineage with respect to its intercontinental disjunctions, since all members of the section are confined to the Mediterranean region and western Asia.



From Africa, samples for the three Malagasy endemics (*Linum betsiliense* Baker, *L. emirnense* Bojer, and *L. marojejyense* (Humbert) Rogers; Rogers 1981b) and the central African *L. keniense*, were not available due to the rarity and age of collections. Available specimens of two South African species, *L. brevistylum* Rogers, and *L. pungens* Planch., are not represented due to recalcitrant DNA extracts and limited available specimens.

From South America, sequences from seven of the 18 accepted species could not be obtained: the two Galapagos endemics (*L. harlingii* Elias. and *L. cratericola* Elias.), the Ecuadorean *L. filiforme* Urb. and Peruvian *L. polygaloides* Planch. (*L. oligophyllum* morphological group according to Mildner 1971), and three rarely collected taxa native to Brazil (*L. organense* Gardn., *L. smithii* Mildner, and *L. palustre* Gardn.).

From North America, four species of *Linum* were not sampled: *L. gypsogenium* G.L. Nesom, *L. modestum* C.M. Rogers, and *L. mcvaughii* C.M. Rogers are all narrow endemics of Mexico, and *L. macrocarpum* C.M. Rogers is known only from the type collection (from southern Alabama). Only four of the 13 species (Sharsmith 1961, O'Donnell 2006) of *Hesperolinon* were sampled.

All of the remaining species of yellow-flowered *Linum* currently known from Africa (Smith 1966, Rogers 1981a), the Americas (Mildner and Rogers 1978, Rogers 1984), and Eurasia (Davis 1967, Ockendon and Walters 1968, Yuzepchuk 1974) were sampled, some with multiple specimens, for a total of 122 samples (90 species, including the outgroup, *Radiola linoides* L.). Accession voucher information and geographic distributions are provided in Appendix B. For divergence time estimation and hypothesis

testing, a reduced dataset was created in order to avoid effects of missing data (since sequences for some markers were not obtainable for some accessions). All accessions with 20% or more of the total sequence length coded as missing data were removed, and most species with multiple accessions were reduced to a single representative to reduce computational time. The *trnK* 3' intron data was not included in the reduced matrix, since sequence from that region could not be obtained from any South American species.

**Topologies used for area cladograms** — Three maximum-likelihood topologies were obtained in the previous phylogenetic study of yellow-flowered *Linum* (Chapter 3) and are referred to here for their biogeographic implications: the combined chloroplast (CPL) marker phylogeny (Figs. 4.3, 4.4), the nuclear ITS phylogeny (Fig. 4.10), and the combined CPL+ITS phylogeny (Fig. 4.12). Details of the data sets and phylogenetic analyses that produced these topologies can be found in Chapter 3. Species are assigned to seven general geographic regions based on distributional ranges as reported in taxonomic and floristic treatments and discerned from natural history collections: Eurasia (EUR), Tropical East Africa (TEA), South Africa (SAF), Eastern South America (eSAM, east of the Andes), Western South America (wSAM, west of the Andes), Southern North America (sNAM, including Mexico and Central America), and North America North of Mexico (nNAM). These geographic ranges are depicted in Fig. 4.1 B. With a few exceptions involving widespread North American species that occur in both MEX and NAM, no species occurs in more than one area. Areas were coded as unordered character states and mapped onto the three maximum likelihood topologies in Mesquite version 2.5

(Maddison and Maddison 2005) using “most-parsimonious reconstructions” mode to elucidate major biogeographic disjunctions among lineages.

**Phylogenetic Analysis of the Reduced Matrix** — The reduced data matrix described above was analyzed using GARLI 0.96b (Zwickl 2006) and Mr. Bayes 3.1.2 (Huelsenbeck and Ronquist 2001), as described in Chapter 3 for the full data matrices. Appropriate models of nucleotide substitution were determined for each partition of the reduced matrix using the Akaike Information Criteria (AIC) implemented in Modeltest 3.7 (Posada and Crandall 1998), so that the appropriate model type could be applied to each data partition during the Bayesian analysis. The phylogeny with the lowest  $-\ln L$  value obtained by GARLI was designated as the observed “unconstrained” topology for comparison to trees found under different constraints during hypothesis testing.

**Hypothesis Testing using the SOWH Test** — The parametric bootstrap procedure termed the SOWH Test (Goldman et al. 2000) was used to test the monophyly or non-monophyly of different lineages, based on the expectations of different biogeographic scenarios, for their suitability as explanations for our data. First, an optimal topology was determined with each hypothetical constraint enforced in GARLI. The likelihood score differences between these constrained topologies and our original unconstrained phylogeny are the test statistics used to determine whether the constraint hypothesis is a possible valid explanation for our data, or, in other words, that the optimal unconstrained topology is a result of chance variation while the constrained topology is the true topology. Substitution model parameters estimated from the observed data given each constrained topology were used to simulate new matrices, each representing a

random result of sequence evolution given the assumption that the constrained topology is true. These simulated matrices were then analyzed with and without the constraint enforced, to obtain a null distribution of score differences that can be expected due to chance variations in sequence evolution. If the test statistic for each constraint is greater than 95% of the null distribution values, then we conclude that it is unlikely ( $p < 0.05$ ) that the constraint is rejected as a potential underlying evolutionary history for the yellow-flowered clade. If the constraint can not be rejected, then we must conclude that the constraint topology is a possible underlying explanation for our observed data, and that our optimal unconstrained topology differs from the constraint merely due to chance. Constraints tested are described as follows:

A: North American Monophyly. The taxa of North America (including Mexico) are constrained to monophyly. If rejected, the implications are that yellow-flowered *Linum* colonized North America at least twice independently, or that a different continental lineage is derived from (nested within) North American ancestry. If not rejected, then we can not rule out a single introduction to North America, and no transfers out of North America.

B: South American non-monophyly. South America is forced to be non-monophyletic in this constraint. If rejected, we conclude that South American monophyly is indeed a significantly better explanation for our data, and that yellow-flowered *Linum* was introduced to South America once. If we can not reject this hypothesis, we must potentially allow for multiple introductions to South America, or South American ancestry for species currently found on another continent.

C: Southern North American Monophyly. The endemic *Linum*s of Mexico and Central America are constrained to monophyly. If rejected, we would conclude either that some Mexican and Central American *Linum*s have distinct origins from other areas and independently colonized the region, or that some other lineages of *Linum* are derived from Mexican or Central American ancestry. If this hypothesis can not be rejected, we must allow for the fact that monophyly of southern North American taxa can't be ruled out, and that these species may share a common origin and are products of speciation within the region.

D: South African non-monophyly. The South African species group is forced to be non-monophyletic under this constraint. If rejected, we can conclude that the South African group is probably monophyletic, that the extant species of yellow-flowered *Linum* in South Africa result from a single colonization event followed by speciation within South Africa, and that no other lineages are derived from within this radiation. If we can not reject this hypothesis as a possible explanation for our data, then we must allow for the possibility that South Africa was colonized multiple times by distinct ancestors, or that at least one South African lineage colonized another continent.

E: New World Non-monophyly. Under this constraint, the New World taxa are forced to be non-monophyletic. If rejected, we would conclude that the New World taxa comprise a single lineage, stemming from a single amphi-Atlantic disjunction event. If we can not reject this constraint, we must allow for the possibility of multiple amphi-Atlantic events, either as multiple introductions to the New World or one introduction to the New World followed by recolonization of the Old World from New World ancestry.

F: Monophyletic South American + South African Clade. All South American and South African Taxa are confined to a single monophyletic group. Rejection would indicate that South American and South African taxa do not share most recent common ancestry. Failure to reject would imply that these continental lineages may be sister to each other, which would be compatible with Gondwanan Vicariance (although existing divergence time estimates for Linaceae already indicate that the diversification of Linaceae significantly post-dates the breakup of Gondwana). This could also potentially be interpreted as an African origin of South American *Linum* by dispersal.

Constraint trees for each hypothesis were generated with MacClade 4.08 (Maddison and Maddison 2005). In the constraint trees, the indicated groups are represented as monophyletic, but no relationships were specified within the constrained clade or among any other taxa in the tree. For the non-monophyly constraints (B, D, and E), the constraint trees (in which the group in question is represented as monophyletic) were enforced as negative constraints (only trees *not* compatible with the constraint tree were allowed), forcing the analysis to find the optimal topology in which the group is *not* monophyletic. For the other hypotheses (A, C, E, and F), the constraint trees were enforced as positive constraints (only topologies compatible with the constraint tree are allowed), resulting in optimal topologies in which the designated group *is* monophyletic. To find the best constrained topology and likelihood score, 100 search replicates were performed in the reduced data matrix in Garli 0.96b with the constraint enforced, and the lowest -LnL score found under each constraint used as the optimal score for that

constraint. The optimal unconstrained score was subtracted from these values to obtain the test statistic for each constraint.

The GARLI-optimized GTR+ $\Gamma$ +I substitution model parameters obtained during each constrained analysis of the original data were used to generate the simulated data matrices on each resulting optimal constrained topology. One-hundred datasets were simulated on each constrained topology using Seq-Gen v. 1.2.7 (Rambaut and Grassly 1997), operated via the *SG Runner* interface (Wilcox 2005). To control for any effects of alignment gaps on the optimal topologies or likelihood scores, the pattern of missing data and gaps found in our original data alignment was replicated in the simulated matrices using a simple Objective-C program written for Mac OS-X by Marc Caddell, which scans the sequence of each taxon on the original matrix for cells coded as missing or gap characters, then replaces any A, C, T, or G at the same sequence position in each simulated matrix with a missing/gap symbol. Each simulated matrix was analyzed with and without its constraint hypothesis enforced in GARLI 0.96b to obtain the null distribution of tree score differences for each constraint.

**Divergence Time Estimates** — A likelihood ratio test was used to test for rate constancy in the combined reduced data matrix; the hypothesis of a molecular clock was rejected ( $P < 0.001$ ). To assess the compatibility of lineage divergence times with potential migration pathways such as the North Atlantic land bridge, divergence times were estimated using BEAST version 1.4.8 (Drummond and Rambaut 2007), which estimates divergence times and confidence intervals using a Bayesian methodology that is not conditional on a single tree topology or set of branch lengths. This is advantageous

in the case of the yellow-flowered clade, since several important nodes in the tree backbone lack significant support in our phylogenetic analyses. The uncorrelated lognormal relaxed clock option was used in BEAST, with a Yule process model for speciation rate. The only prior set to a non-default value was the *tmrca* prior – the age of the root of the tree was set to a normal distribution prior with the mean at 28.5 mya before present and a standard deviation of 1.0. This is the average divergence time estimated in Chapter 2 for the split of *Radiola linoides* from yellow-flowered *Linum*, in analyses that included a fossil calibration point at 33.9 my for the minimum age of the common ancestor of all *Linum* sp., based on the pollen fossil described by Cavagnetto and Anadon (1996), and constrained the maximum age of Linaceae to values estimated by Davis et al. (2005). The MCMC chain was run for 30 million generations in BEAST with the auto-optimize feature activated (burn-in period of 3 million generations). Parameters were sampled every 1000<sup>th</sup> generation. The run was replicated to verify post-burn-in convergence of on similar likelihood values and divergence time estimates, and results from the two runs combined as recommended in the BEAST manual. Effective sample sizes for all estimated parameters and node ages were well above 200 (also as recommended).

## RESULTS

The ITS (Fig. 4.3) and CPL (Fig. 4.4) topologies agree in several respects: the basal position of Eurasian taxa (indicating Eurasian ancestry for the African and New World lineages), the monophyly of South African taxa (indicating a single introduction to



that continent), and the sister relationship of Tropical African linums (represented by *Linum volkensii* and *L. holstii*) to *L. strictum* (a Mediterranean species). They differ, however, in terms of the biogeographic relationships among African and New World species groups. In the ITS topology, the South African clade is shown as sister to a North American clade consisting of *Hesperolinon*, *Sclerolinon*, and a group of Mexican endemics that has been referred to as the *Linum mexicanum* group. The ITS topology (Fig. 4.3) also shows a Mexican species, *Linum longipes*, inserted among the South American taxa. A Mexican/Central American distribution is indicated for the ancestor of the South African, South American, and northern North American lineages, with South Africa being colonized once and both South and northern North America colonized twice. The ITS data does not provide strong support for any of the most important nodes along the backbone of the phylogeny, however. The CPL topology (Fig. 4.4) shows the New World lineages as monophyletic, with northern North America as the first New World area colonized. The CPL topology implies that South America was colonized once from northern ancestry, and Mexico perhaps multiple times, with possible recolonizations of northern North America from Central American ancestry.

The combined unreduced topology (Fig. 4.5) again shows a sister relationship between the tropical African taxa and Eurasian *Linum strictum*. The node representing the most recent common ancestor of the South African and New World linums is equivocal; this ancestor could have resided on any of the continents in question. The South African species complex is monophyletic and well supported, and the New World lineage is moderately well supported as monophyletic. The New World taxa are divided

among three clades, but the relationships among these clades is not well supported or resolved. The South American species complex is well supported as monophyletic and derived from North American ancestry, but its specific origin from either Mexican or more northern ancestry is uncertain. The North American taxa are divided into two lineages, one including the North American segregate genera and the *Linum mexicanum* group, and the other with the majority of North American species. The affinities of the Mexican endemic *Linum longipes* are not well resolved.

**Reduced data unconstrained phylogeny** — The final reduced matrix consisted of 74 accessions representing 72 species (including the outgroup, *Radiola linoides*), and 3086 bp of combined chloroplast and nuclear data (1160 bp matK, 440 ndhF, 837 trnL-F, 649 ITS). Accessions included in the reduced matrix are indicated in Appendix B. Data matrix characteristics for the reduced data set are given in Table 4.1. The phylogeny resulting from this matrix is presented in Fig. 4.6, with geographic areas traced on the phylogeny as described above. This phylogeny is largely compatible with those derived from the original taxon sample in Chapter 3. The Old World Northern Hemisphere species of *Linum* are found to be monophyletic, and sister to a well-supported South African and New World clade. South African taxa are a well-supported group sister to a New World lineage. The South American lineage is well supported as monophyletic, and appears to be inserted between two North American lineages, but the branching order of these lineages is not well supported. The monophyly of each North American clade is not well-supported, with considerable uncertainty indicated for the relationships of *L.*

*longipes*, the series *Virginiana* clade (*L. westii* to *L. medium*), and the *L. neomexicanum*-*L. kingii* clade.

**Parametric Bootstrap Results** — The optimal unconstrained phylogeny had a -LnL score of 15354.06. Optimal likelihood scores resulting from constrained analyses are shown in Table 4.2, along with SOWH test results for each hypothesis. All constraint hypotheses were rejected with  $P < 0.01$  except for that of New World non-monophyly ( $P = 0.12$ ).

**Divergence Time Estimates** — Mean divergence times and 95% confidence intervals ( = 95% height density proportion) estimated in BEAST for key nodes of the yellow-flowered *Linum* phylogeny (nodes designated by letters in Fig. 4.5) are given in Table 4.3. Divergence times and confidence intervals are presented graphically in Fig. 4.7; node heights (distance from each node to the root) in this chronogram represent the average divergence time estimated by BEAST during the post-burn-in sampling. Most of the major lineages of yellow-flowered *Linum* are estimated to have originated in the Miocene, though the South African and New World lineage may have diverged from its Old World Northern Hemisphere relatives in the early Oligocene (32.22 mya, maximum). Most of the species sampled appear to have diverged from their sister species in the Pliocene (Fig. 4.7). Divergences of the South African, South American, and North America lineages from each other (nodes E, F, and H) are estimated to have occurred within a narrow span of time in the mid-Miocene, between 16 and 15 mya, although confidence interval indicate these divergences may have occurred any time between 26.54 and 9.5 mya. Extant sampled South African species are estimated to have radiated

from their common ancestor relatively recently, between 4 and 9 mya (avg. 6.21), compared to North American and South American lineages.

## DISCUSSION

We find that a single trans-Atlantic event provides the most parsimonious explanation for the amphi-Atlantic disjunction in the yellow-flowered lineage of *Linum*, indicated by the monophyly of the New World taxa in the optimal phylogenies resulting from all unconstrained analyses of the combined ITS and CPL data (Figs. 4.4, 4.5). However, considerable uncertainty surrounds this issue. First, the branching order of lineages along the “backbone” of the phylogeny of yellow-flowered *Linum* lacks significant posterior probability or bootstrap support, due either to conflicting or inadequate phylogenetic signal from the combined markers. The ITS data alone suggest that the South African clade is nested within the “New World” lineage, sister to the North American *Hesperolinon* and *Linum mexicanum* group clade, which potentially requires a second trans-Atlantic event, from North America to South Africa (Fig 4.3). The parametric bootstrap test of New World monophyly subsequently fails to reject this hypothesis, so we can not rule out the possibility of two trans-Atlantic events, either as two colonization’s of the New World or the colonization of South Africa from New World ancestry. The optimal resolutions for the combined data matrices, which place the South African clade sister to the New World clade, also create a situation in which it is not possible to ascertain (via parsimony area optimizations) which continent, Africa or

Eurasia, was the origin of the New World colonist(s). Rejection of South African non-monophyly argues against a South African origin for the New World clade.

Divergence time estimates indicate that New World yellow-flowered *Linum* diverged from extant Old World taxa and began to diversify in the Americas during the Miocene, between 22.5 and 10 mya (Fig. 4.7). This is compatible with migration across the remnants of the North Atlantic land bridge during the last stages of its existence (Milne 2006). Several other angiosperm lineages exhibit Miocene amphi-Atlantic disjunctions that are concluded to be products of North Atlantic migration, including Cistaceae (Guzmán and Vargas 2009), *Cercis* (Davis et al. 2002b) *Platanus* (Meyers and Liston 2008), *Valerianella* (Bell 2007), *Fagonia* (Beier et al. 2004), Burseraceae (Weeks and Simpson 2005), and multiple lineages of Antirrhineae (Vargas et al. 2004). Like *Linum*, several of these are taxa whose seeds and fruits are thought to have very limited potential for long-distance dispersal (eg., *Cercis*, *Fagonia*, and Antirrhineae, all of which were noted bear dehiscent capsules and relatively large, smooth, unwinged seeds).

Two independent amphitropical disjunctions are indicated by our results. In the New World, parsimony ancestral area reconstructions on all optimal topologies indicate that South American *Linum* is derived from North American ancestry (Figs. 4.3, 4.4, 4.5, 4.6). We interpret the rejection of North American monophyly as an argument in favor of the North American origin of South American *Linum*. South American *Linum* is concluded to be monophyletic (non-monophyly rejected by SOWH test,  $P < 0.01$ ), and therefore a result of a single introduction to the continent followed by speciation. The colonization of South America is estimated to have occurred in the latter part of the

Miocene, no later than 6 mya (node P, Fig. 4.7, the time estimated for the beginning of yellow-flowered diversification in South America). During this period, the closure of the isthmus of Panama is thought to have been in its final stages (Keigwin 1978), with potential volcanic island arcs providing a possible migration pathway up until completion of the isthmus in the Pliocene. In spite of its limited dispersal ability, *Linum* might therefore be added to the list of taxa providing evidence for dispersal between North and South America prior to closure of the isthmus (Ickert-Bond et al. 2009). *Linum* appears to have diversified first in Eastern South America, with several lineages colonizing the western portion of the continent as *Linum* diversified in late Miocene and Pliocene.

The South African species group is generally well supported as monophyletic (Figs. 4.3-4.6), implying a single introduction of yellow-flowered *Linum* from the Northern Hemisphere to South Africa, followed by speciation within South Africa. South African non-monophyly is rejected ( $P < 0.01$ ), further supporting the conclusion that *Linum* was introduced to South Africa once. With the exception of the widespread species *Linum thunbergii*, the most widespread species of *Linum* in Africa which we conclude originated in the South African radiation and then spread northward into Tropical East Africa, no dispersals or migrations out of South Africa are apparent (Figs. 4.3-4.7). The hypothesis that the Tropical African taxa provide a biogeographic connection between Eurasia and South Africa, as proposed by Rogers and observed for some South African clades in other families (e.g. Penaeaceae and Stilbaceae; Galley and Linder 2006), is refuted by our results. The tropical African endemic taxa sampled (*Linum holstii* and *L. volkensii*) are clearly derived independently from Eurasian ancestry,

and do not share a common origin with the South African group. South African *Linum* are estimated to have diverged from the other continental lineages some 16 million years ago (node E, Fig. 4.7), with the sampled extant species beginning their diversification in South Africa between 3.5 and 9 mya (node W, Table 4.3, Fig. 4.7). This dating makes *Linum* a relatively recent addition to the South African Cape flora compared to some other Cape lineages whose divergence times from non-Cape relatives have been estimated to the mid-Oligocene or earlier (Galley and Linder 2006). Though the sister group of the South African lineage of *Linum* is apparently a New World clade, the exact origin of the South African lineage, whether Eurasia or North America, can not be determined from our current data and taxon sampling. Additional sampling from the native flaxes of Madagascar and India, areas often implicated in the biogeographic histories of the South African Cape Flora (Galley and Linder 2006), may clarify the origins of South African *Linum* in the future.

## CONCLUSIONS

While the relationships of North American, South American, and South African species groups are poorly resolved by our current data, the parametric bootstrap method used here allows some conclusions to be made about the biogeographic history of yellow-flowered *Linum*. Both South America and South Africa appear to have been colonized only once — South America from North America, and South Africa from either Eurasia or North America. The intercontinental disjunct lineages of yellow-flowered *Linum* are estimated to have originated in the Miocene, when land bridges and island arcs may have

facilitated migration between Eurasia and North America and from North America to South America. In accordance with the probable low dispersability of *Linum* seeds and fruits, we find no compelling evidence for recent intercontinental long-distance dispersal in *Linum*; dispersal is generally invoked only for more recent divergences, usually of Pliocene age (Wen and Ickert-Bond 2009). Additional data and sampling are certain to resolve some of the relationships more confidently, and will allow better estimates of divergence times (Sanderson 1998).



Table 4.1: Data matrix characteristics for the reduced taxon set used for divergence time estimation and biogeographic hypothesis testing. NTax = number of terminal taxa (ndhF sequence missing for *L. quadrifolium*), Nchar = total number of base pairs in aligned matrix for each region. Constant = # of constant characters. P.i. = number of parsimony informative characters. Model=substitution model type selected.

Matrix:	<i>matK</i>	<i>ndhF</i>	<i>trnL-F</i>	CPL Combined	ITS	CPL + ITS
NTax	74	73	74	74	74	74
NChar	1160	440	837	2437	649	3086
Number included	1160	440	837	2437	649	3086
constant	900	363	654	1918	340	2258
p.i.	132	41	93	266	228	494
Model	GTR+ $\Gamma$ +I	GTR+ $\Gamma$ +I	GTR+ $\Gamma$ +I	GTR+ $\Gamma$ +I	GTR+ $\Gamma$	GTR+ $\Gamma$ +I

Table 4.2: SOWH test results for six constraints imposed on the reduced molecular dataset.

Constraint Hypothesis	-LnL	$\Delta$	P
unconstrained	15354.06		
A) North American Monophyly	15360.43	6.37	<0.01
B) South American non-Monophyly	15365.64	11.58	<0.01
C) S. North American Monophyly	15487.82	133.76	<0.01
D) South Africa non-Monophyly	15391.81	37.75	<0.01
E) New World non-monophyly	15354.88	0.8282	0.12
F) South America+Africa monophyly	15363.96	9.90	<0.01

Table 4.3: Divergence times (mya) for lineages of yellow-flowered *Linum*, estimated by BEAST. Letters correspond to nodes in Figs. 4.6 and 4.7.

MRCA of...	mean	geom. mean	median	95% confidence interval	
				lower	upper
(root) Radiola and yellow-flowered clade	28.07	28.00	28.06	24.20	32.10
A) Yellow-flowered <i>Linum</i> and segregate genera	26.35	26.06	27.13	17.82	32.22
B) Old World Northern Hemisphere group	24.55	24.11	25.37	15.36	31.52
C) <i>Cathartolinum</i> , <i>Syllinum</i> , <i>Dichrolinum</i> clade	22.64	22.14	22.84	13.68	30.73
D) MRCA of sect. <i>Syllinum</i> and subsect. <i>Halolinum</i>	18.21	17.73	18.10	10.12	26.54
E) South African and New World <i>Linum</i>	16.53	16.27	16.48	10.70	22.36
F) New World <i>Linum</i>	16.13	15.87	16.09	10.46	21.78
G) <i>Hesperolinon</i> , <i>Sclerolinon</i> , <i>L. mexicanum</i> group	15.14	14.83	15.00	9.36	21.12
H) MRCA of South American and main North American clade	14.97	14.73	14.93	9.54	20.12
I) main NAM clade	14.60	14.36	14.54	9.43	19.79
J) <i>Halolinum</i> , OWNH subsect. <i>Linopsis</i> clade	14.49	14.07	14.33	7.72	20.85
K) MRCA of <i>L. kingii</i> and <i>L. carteri</i>	14.06	13.82	14.01	8.98	19.06
L) MRCA of <i>L. schiedeanum</i> and <i>L. carteri</i>	12.37	12.15	12.33	7.71	16.68
M) MRCA <i>L. strictum</i> and <i>L. volkensii</i>	11.81	11.39	11.60	6.08	17.92
N) <i>Hesperolinon</i> and <i>Sclerolinon</i>	11.38	11.13	11.27	6.93	16.09
O) subsect. <i>Rigida</i> and <i>L. rupestre</i> group	10.12	9.93	10.07	6.35	13.93
P) South American clade	9.93	9.73	9.85	5.96	13.90
Q) sect. <i>Linopsis</i> subsect. <i>Halolinum</i> )	9.64	9.28	9.45	4.77	14.76
R) <i>Linum schiedeanum</i> group	9.24	9.01	9.16	5.19	13.25
S) Subsection <i>Rigida</i>	8.64	8.47	8.59	5.41	11.98
T) MRCA of wSAM <i>Linum</i>	7.91	7.73	7.81	4.61	11.20
U) <i>Hesperolinon</i>	7.47	7.25	7.34	4.05	11.04
V) Series <i>Neo-Mexicana</i>	6.29	6.00	6.11	2.70	10.00
W) South African <i>Linum</i>	6.21	6.04	6.10	3.49	9.06
X) <i>Virginiana</i> clade	6.15	5.90	5.94	3.06	9.73
Y) <i>Linum mexicanum</i> group	3.08	2.85	2.86	1.09	5.64
Z) section <i>Syllinum</i>	2.53	2.36	2.38	0.92	4.41
A1) <i>Linum rupestre</i> group	2.04	1.89	1.90	0.71	3.70

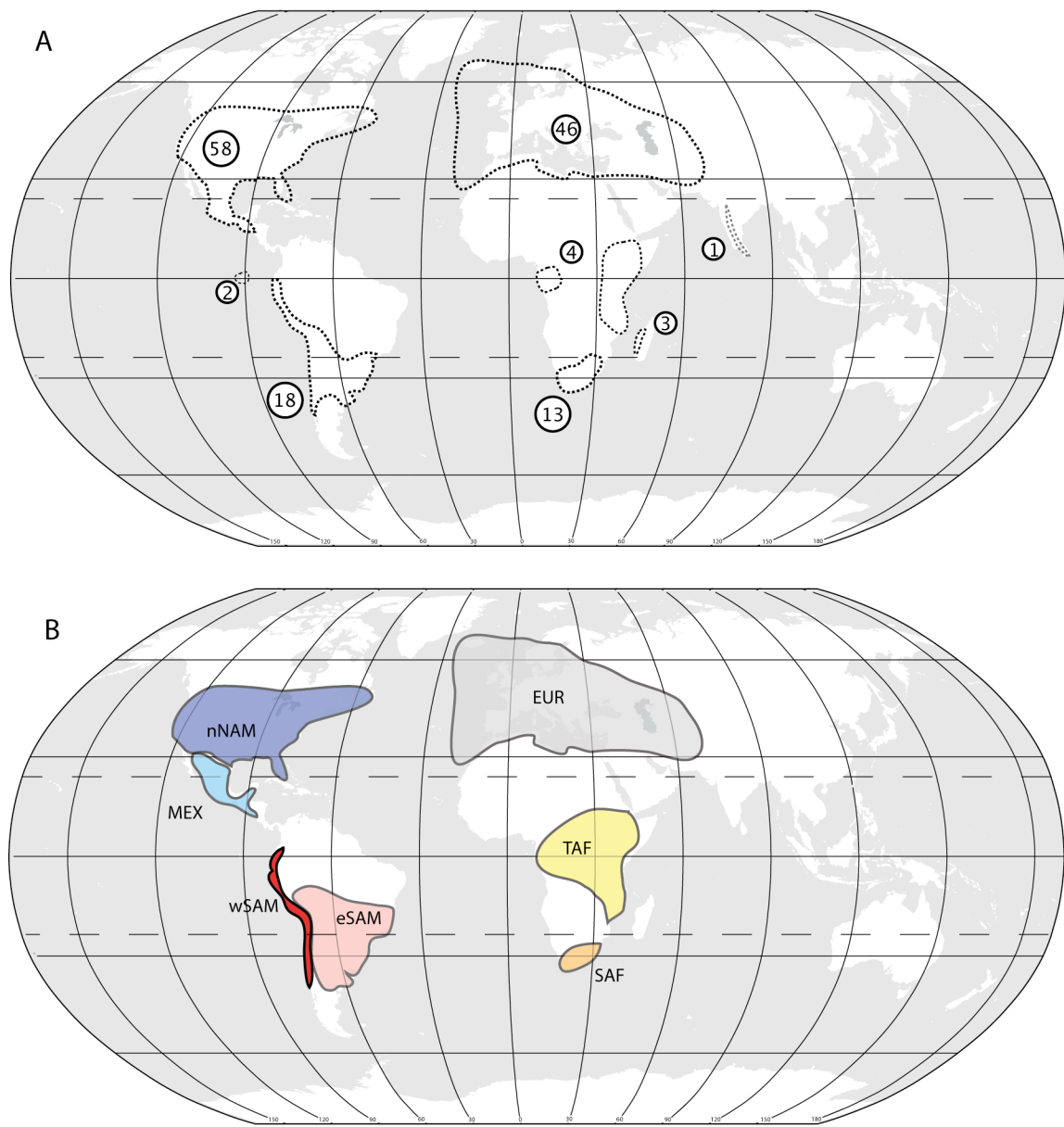


Fig 4.1. Distribution maps for yellow-flowered clade. A) Composite distribution of the “yellow-flowered” flaxes (including *Linum* sections *Linopsis*, *Syllinum*, and *Cathartolinum*, and the segregate genera *Cliococca*, *Hesperolinon*, *Radiola*, and *Sclerolinon*), and approximate number of species in each area. B) Major geographic areas to which species included in this study were assigned. EUR= Eurasia, nNAM= North America north of Mexico, MEX=Mexico endemic, wSAM=South America west of the Andes, eSAM=South America east of the Andes, TAF=Tropical/Central/East Africa, SAF=South Africa.

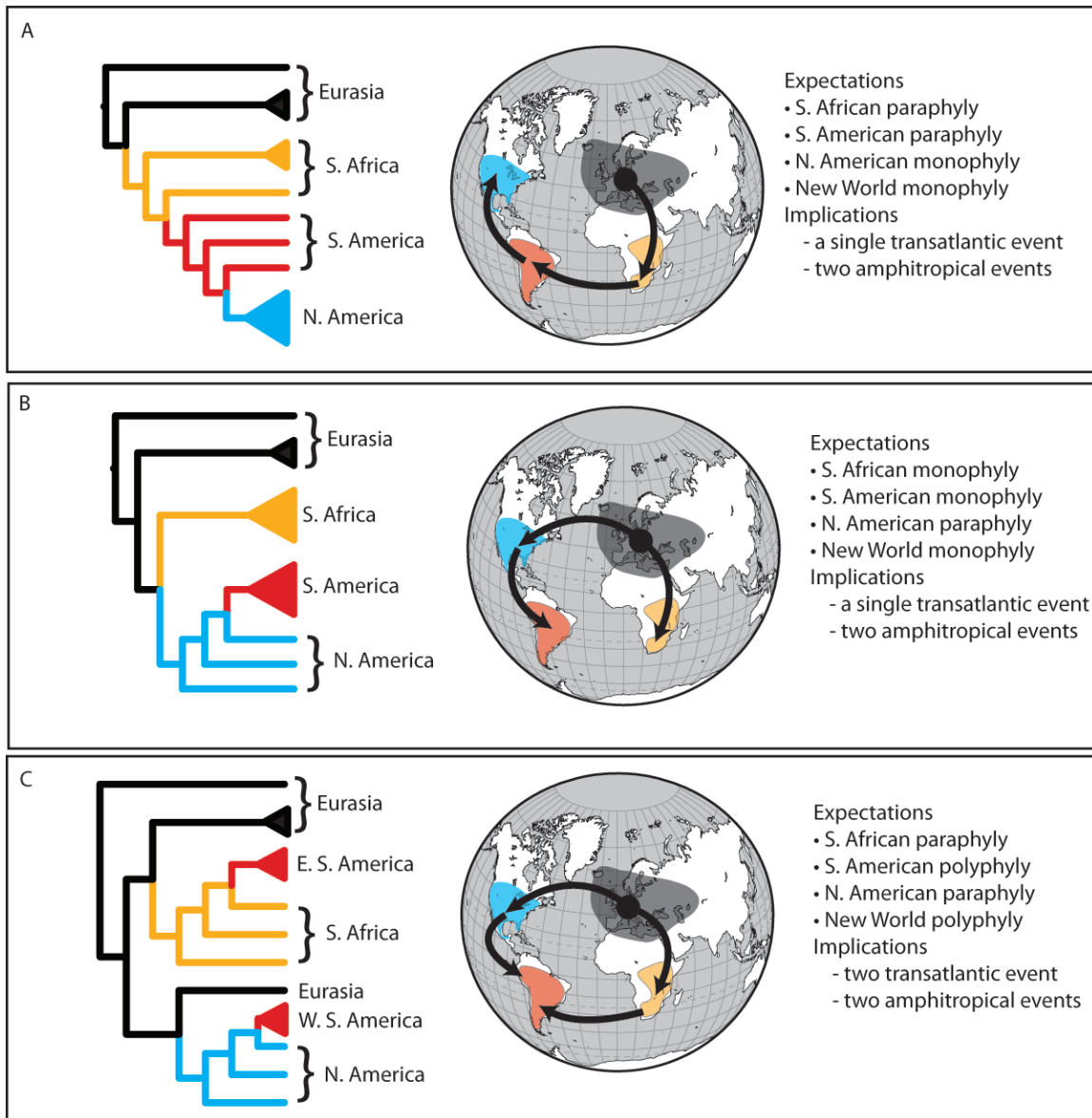


Fig. 4.2. Hypothetical historical biogeographic scenarios for yellow-flowered *Linum*, with simplified phylogenetic expectations. A) A South African origin of New World *Linum* might imply paraphyly for the South African taxa, with American taxa nested among them. B) A single amphi-Atlantic disjunction and independent Old and New world amphitropical events could imply monophyly for both South American and South African groups, and paraphyly of North American linums. C) More complex scenarios involving multiple trans-Atlantic events could result in paraphyly or polyphyly of species groups on multiple continents.



## 128

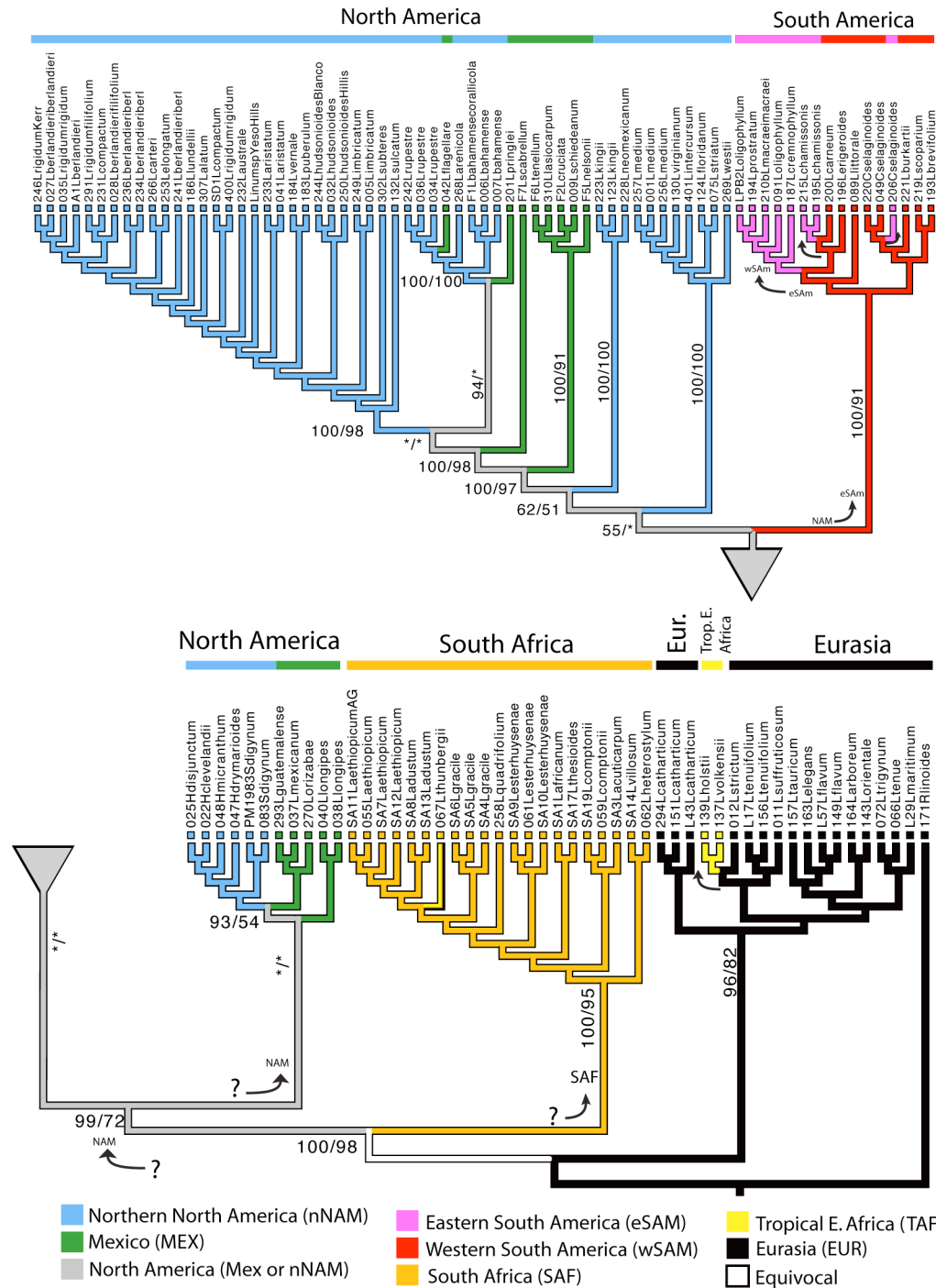


Fig 4.5: Parsimony mapping of biogeographic areas on combined (unreduced) CPL+ITS molecular topology. Bayesian posterior probabilities (as %) and ML bootstrap % above branches (for relevant nodes only). Values below 50% indicated by \*.

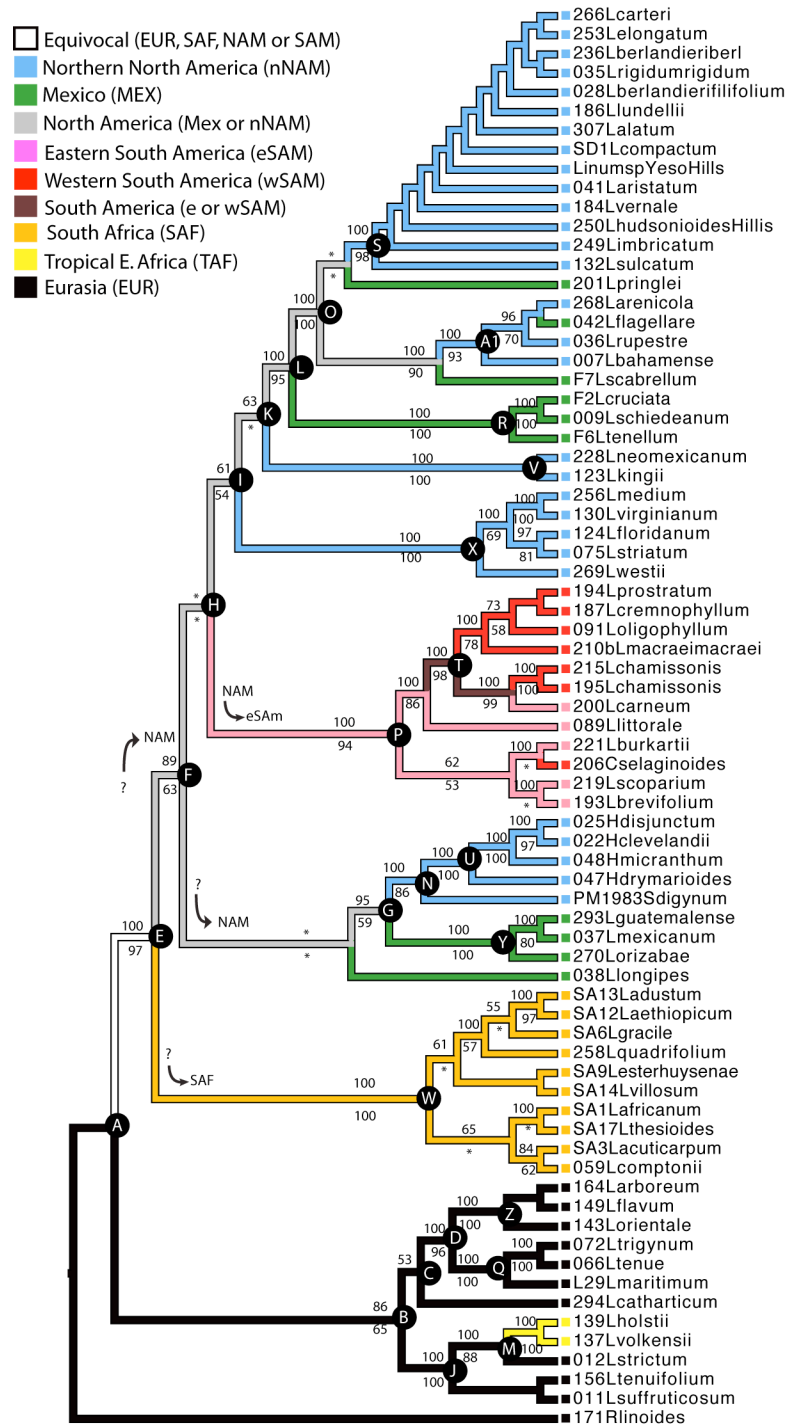


Fig. 4.6. Optimal ML topology for reduced data matrix, with parsimony mapping of geographic areas. Bayesian PP (as %) above branches, ML bootstrap below (<50% indicated by \*). Lettered nodes correspond to nodes for which divergence times were estimated in BEAST, indicated in Table 4.3 and Fig. 4.6.



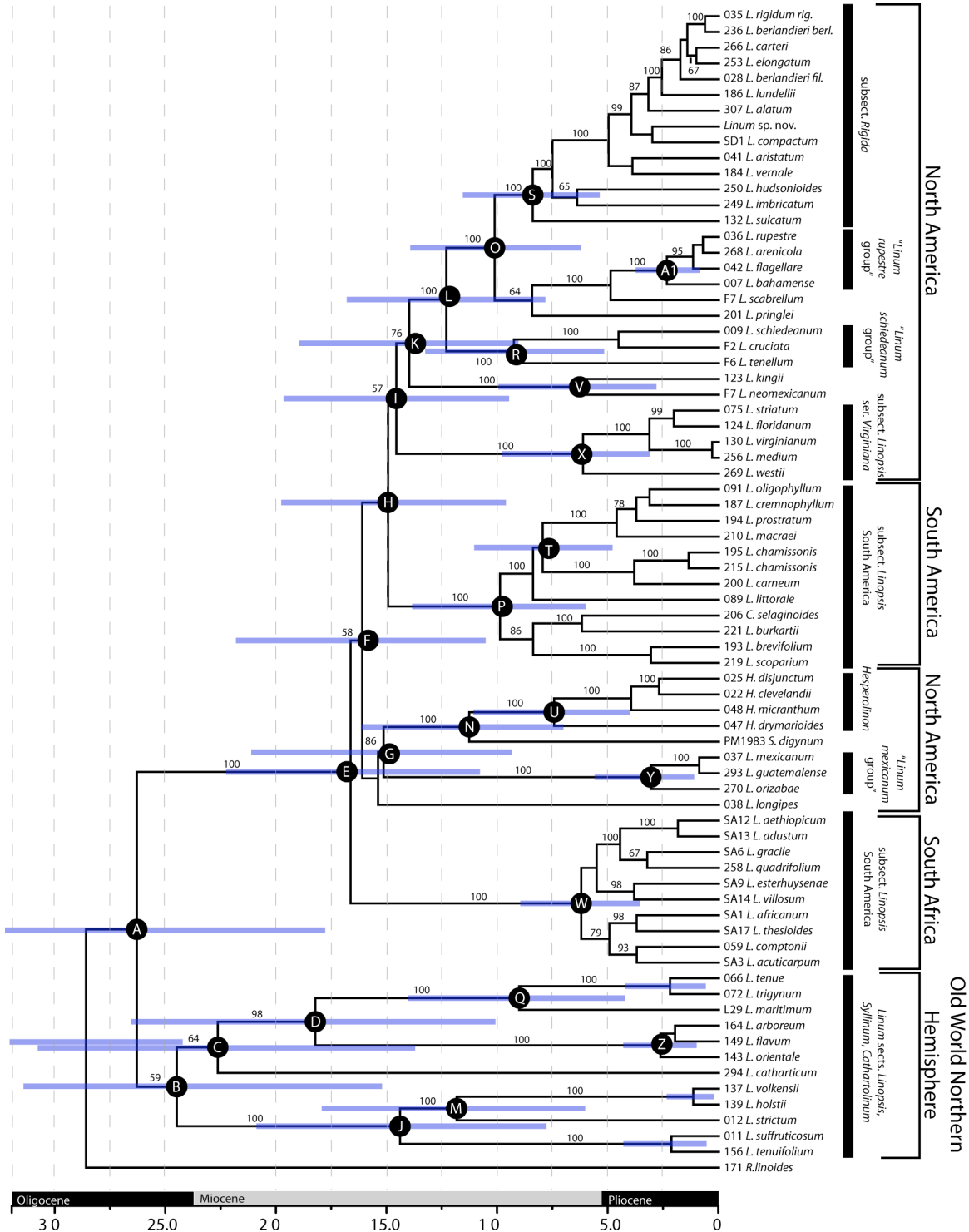


Fig. 4.7. Majority-rule consensus of trees sampled during BEAST analysis, rooted with *R. linoides*, with node heights adjusted to mean divergence times. Posterior probabilities above branches (as %, values < 50 not shown). Bars indicate 95% confidence interval around mean divergence time for each node. Letters correspond to clades in Fig. 4.5 and Table 4.3.

## Appendix A: *rbcL* and *matK* alignment, Chapter 2

*rbcL* (1-1293), *matK* (1294-.)

Taxon/Node	111111111122222222223333333333444444444455555555556666666666777
	12345678901234567890123456789012345678901234567890123456789012
L. lewisii	TATAAATTAACCTTATTATACTCCTGAATATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. perenne	TATAAATTAACCTTATTATACTCCTGAATATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. pratense	TATAAATTAACCTTATTATACTCCTGAATATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. narbonense	TATAAATTAACCTTATTATACTCCTGAATATGAAACCAAAGATACTGATATCTTGGCAGCATTC
C. selaginoides	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. hirsutum	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. hypericifolium	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. usitatissimum	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. nervosum	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. bienne	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. monogynum	----AATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. suffruticosum	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. clevelandii	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. disjunctum	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. berlandieri	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. rupestre	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
D. pentagyna	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
R. indica	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. drymarioides	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. micranthum	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. castaneifolia	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
S. digynum	-----AATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. oligophyllum	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. neblinae	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. mystax	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC	
H. gabunensis	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. obtusifolia	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. neocaledonic	-----AGTAACCT
H. penicillanth	-----CCAAAGCTACTGATATCTTGGCAGCA-----AGTAACCT
H. spicata	-ATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. planchoni	---AATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. orientalis	----AATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. longipes	-ATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. volkensii	TATAAATTAACCTTATTATACTCCTGAATATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. flavum	TATAAATTAACCTTATTATACTCCTGAATATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. catharticum	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. arboreum	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
R. linoides	TATAAATTAACCTTATTATACTCCGAATATGAAACCAAAGATACTGATATCTTGGCAGCATTC
R. schomburgkii	---AATTGACTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
R. monsalveae	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
A. pubescens	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
A. khasayama	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
A. saxatilis	--TAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
L. comptonii	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
H. humiriifolia	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
I. griffithiana	-----
P. magnifolia	-----
T. sinensis	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
T. ovoidea	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
R. calophylla	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
AcalyphaInsulana	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
BalanopsPancheri	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
BischofiaJavanic	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
BruguieraGymnorh	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
ByrsonimaCrassif	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
MalpighiaGlabra	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC
CamptostylusMann	TATAAATTAACCTTATTATACTCCTGACTATGAAACCAAAGATACTGATATCTTGGCAGCATTC



H. neocaledonic	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
H. penicillanth	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
H. spicata	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
H. planchoni	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
H. orientalis	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
H. longipes	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
L. volkensii	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
L. flavum	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
L. catharticum	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
L. arboreum	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
R. linoides	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
R. schomburgkii	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
R. monsalveae	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
A. pubescens	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
A. khasayama	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
A. saxatilis	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
L. comptonii	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
H. humiriifolia	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
I. griffithiana	-----
P. magnifolia	-----
T. sinensis	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
T. ovoidea	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
R. calophylla	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
AcalyphaInsulana	CCTCAACCTGGAGTTCGCGCGGAGGAAGCAGGAGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
BalanopsPancheri	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
BischofiaJavanic	CCTCAACCTGGAGTTCACCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
BruguieraGymnorh	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
ByrsonimaCrassif	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGAGTTCGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
MalpighiaGlabra	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGAGTTCGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
CamplostylusMann	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
ClusiaRosea	CCTCAACCTGGAGTTCACCTGAGGAAGCAGGAGCTGCAGTAGCTGCGAGAATCTTCTACAGGTACATGGACA
CratoxylumCochin	CCTCAACCTGGAGTTCACCTGAGGAAGCAGGAGCTGCAGTAGCTGCGGAATCTTCTACTGGTACATGGACA
CtenolophonEngle	CCTCAACCTGGAGTTCGCGCTGAAGAAGCAGGGGCTGCAGTAGCTGCTGAATCTTCTACTGGTACATGGACA
DrypetesLittoral	CCTCAACCAGGAGTTCGCGCTGAGGAAGCTGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
ElatineTriandra	CCTCAACCTGGGTTCCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
ErythroxylumNovo	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
EuonymusHamilton	CCTCAACCTGGAGTTCGCGCTGAAGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
EuphorbiaHumifus	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGAGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
EuphoniaGuianen	CCTCAACCAGGAGTTCACCTGAAGAAGCAGGGGCTGCGGTAGCTGCGGAATCTTCTACCGGTACATGGACA
FlacourtiaIndica	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
FlueggeaVirosa	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
GarciniaSubellip	CCTCAACCTGGAGTTCACCCGAGGAAGCAGGAGCTGCAGTAGCTGCGGAATCTTCTACAGGTACATGGACA
GomphiaSerrata	CCTCAACCTGGAGTTCGCGCGGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
HumiriaBalsamife	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
IonidiumCommune	CCTCAACCAGGAGTTCGCGCGGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
IringiaMalayana	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
IxonanthesReticu	CCTCAACCTGGTGTTCGCGCGGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
JatrophaIntegerr	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGAGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
KiggelariaAfrica	CCTCAACCTGGAGTTCGCGCTGAAGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
LacistemaAggrega	CCTCAACCTGGAGTTCACCCGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
LunaniaParviflor	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGAGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
ManihotEsculenta	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
OxalisCorniculat	CCTCAACCTGGAGTTCCTCTGAGGAAGCTGGGGCAGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
PassifloraQuadra	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGAGCTGCAGTAGCTGCTGAATCTTCTACTGGTACATGGACA
PhyllanthusFlexu	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
PimelodendronGri	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGAGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
RicinusCommunis	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGAGCTGCAGTAGCTGCTGAATCTTCTACTGGTACATGGACA
SacoglottisSp	CCTCAACCTGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
ScageaOligostemo	CCTCAACCAGGAGTGCCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCGGAATCTTCTACTGGTACATGGACA
TrigoníaBolivian	CCTCAACCAGGAGTTCGCGCTGAGGAAGCAGGGGCTGCGGTAGCTGCGGAATCTTCTACTGGTACATGGACA
AlbiziaTomenosaJ	CCTCAACCTGGAGTTCGCGCTGAAGAAGCAGGTGCGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
KerriaJaponica	CCCCAACCTGGAGTTCACCTGAGGAAGCAGGTGCGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
DatiscaCannabina	CCTCAACCTGGAGTTCACCTGAGGAAGCAGGTGCGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA
QuercusRubra	CCTCAACCTGGAGTTCGCGCGGAGGAAGCAGGGGCTGCGGTAGCTGCTGAATCTTCTACTGGTACATGGACA

	4444455555555566666666667777777777888888888899999999990000000001111111
Taxon/Node	567890123456789012345678901234567890123456789012345678901234567890123456
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L. lewisii	ACCGTGTGGACCGACGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
L. perenne	ACCGTGTGGACCGACGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
L. pratense	ACCGTGTGGACCGACGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
L. narbonense	ACTGTGTGGACCGACGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
C. selaginoides	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
L. hirsutum	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
L. hypericifoliu	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
L. usitatissimum	ACTGTGTGGACCGACGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
L. nervosum	ACCGTGTGGACCGACGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
L. bienne	ACTGTGTGGACCGACGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
L. monogynum	ACTGTGTGGACCGACGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
L. suffruticosum	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
H. cleavelandii	ACTGTGTGGACCGATGGACTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
H. disjunctum	ACTGTGTGGACCGATGGACTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
L. berlandieri	ACTGTGTGGACCGATGGGCTTACTAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
L. rupestre	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
D. pentagyna	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
R. indica	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
H. drymarioides	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
H. micranthum	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
H. castraneifolia	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
S. digynum	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
L. oligophyllum	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
H. neblinae	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
H. mystax	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
H. busseana	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
H. gabunensis	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
H. obtusifolia	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACTACATTGAACCCGTTGCC
H. neocaledonic	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
H. penicillanth	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
H. spicata	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACGGCATTGAACCCGTTGCC
H. planchoni	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
H. orientalis	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
H. longipes	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
L. volkensii	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
L. flavum	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
L. catharticum	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
L. arboreum	ACTGTGTGGACCGATGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
R. linoides	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
R. schomburgkii	ACTGTGTGGACCGATGGACTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
R. monsalveae	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
A. pubescens	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
A. khasayama	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
A. saxatilis	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
L. comptonii	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTCCCT
H. humiriifolia	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
I. griffithiana	-----
P. magnifolia	-----
T. sinensis	ACTGTGTGGACCGACGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
T. ovoidea	ACTGTGTGGACCGACGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
R. calophylla	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATTGAACCCGTTGCC
AcalyphaInsulana	ACTGTGTGGACCGATGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACGACATCGAGCCGTTGCT
BalanopsPancheri	ACTGTGTGGACCGATGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATCGAGCCGTTGCT
BischofiaJavanic	ACCGTGTGGACCGACGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATCGAGCCGTTGCT
BruguieraGymnorh	ACCGTGTGGACCGATGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATCGAGCCGTTGCT
ByrsonimaCrassif	ACTGTGTGGACCGATGGGCTTACCAGCCTTGATCGTTATAAAGGACGATGCTACCACATCGAGCCGTTGCT
MalpighiaGlabra	GCTGTGTGGACCGATGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATCGAGCCGTTGCT
CampostylusMann	AGTGTGTGGACCGATGGACTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACGACATCGAGCCGTTGCT
ClusiaRosea	ACTGTGTGGACCGATGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATCGAGCCGTTGCT
CratoxylumCochin	ACTGTGTGGACCGATGGACTGACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATTGAGCCTGTTCCCT
CtenolophonEngle	ACTGTGTGGACCGACGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTATCACATCGAGCCGTTGCT
DrypetesLittoral	ACTGTGTGGACCGATGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATCGAGCCGTTGCT
ElatineTriandra	ACTGTGTGGACCGATGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTATCACATCGAGCCGTTGCT
ErythroxylumNovo	ACTGTGTGGACCGATGGGCTGACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATCGAGCCGTTGCT
EuonymusHamilton	ACTGTGTGGACCGATGGGCTTACCAGCTCTTGATCGTTATAAAGGACGATGCTACCACATCGAGCCGTTGCT





L. narbonense ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTTAAAGCCCTGCGAGCCCTACGTCTGGAGGATTTGCGA  
 C. selaginoides ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 L. hirsutum ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTTAAAGCCCTGCGAGCCCTACGTCTGGAGGATTTGCGA  
 L. hypericifolium ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTTAAAGCCCTGCGAGCCCTACGTCTGGAGGATTTGCGA  
 L. usitatissimum ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTGCGAGCCCTACGTCTGGAGGATTTGCGA  
 L. nervosum ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTTAAAGCCCTGCGAGCCCTACGTCTGGAGGATTTGCGA  
 L. bienne ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTTAAAGCCCTGCGAGCCCTACGTCTGGAGGATTTGCGA  
 L. monogynum ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTTAAAGCCCTGCGAGCCCTACGTCTGGAGGATTTGCGA  
 L. suffruticosum ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 H. clevelandii ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 H. disjunctum ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 L. berlandieri ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGTCTGGAGGATTTGCGA  
 L. rupestre ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 D. pentagyna ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 R. indica ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTGCGYGCCCTACGCCTGGAGGATTTGCGA  
 H. drymarioides ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 H. micranthum ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 H. castaneifolia ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 S. digynum ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGCCTGGAGGATTTGCGA  
 L. oligophyllum ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 H. neblinae ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 H. mystax ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 H. busseana ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
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 H. obtusifolia ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 H. neocaledonic ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 H. penicillanth ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 H. spicata ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 H. planchoni ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
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 H. longipes ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 L. volkensii ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGTGCCCTACGCCTGGAGGATTTGCGA  
 L. flavum ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGTCTGGAGGATTTGCGA  
 L. catharticum ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 L. arboreum ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGTCTGGAGGATTTGCGA  
 R. linoides ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 R. schomburgkii ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 R. monsalveae ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 A. pubescens ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 A. khasayama ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 A. saxatilis ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 L. comptonii ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 H. humiriifolia ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 I. griffithiana -----  
 P. magnifolia -----  
 T. sinensis ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 T. ovoidea ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCCCTACGCCTGGAGGATTTGCGA  
 R. calophylla ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 AcalyphaInsulana ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCCCTACGTCTGGAGGATTTGCGA  
 BalanopsPancheri ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCTCTACGTCTGGAGGATTTGCGA  
 BischofiaJavanic ATGTTTACTTCCATTGTAGGTAATGTATTTGGGTTCAAAGCCCTACGCGCTCTACGTCTGGAGGATTTGCGA  
 BruguieraGymnorh ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCTCTACGTCTGGAGGATTTGCGA  
 ByrsonimaCrassif ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCTCTACGTCTGGAGGATTTGCGA  
 MalpighiaGlabra ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCTCTCCGTCTGGAGGATTTGCGA  
 CampostylusMann ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCTCTACGTCTGGAGGATTTGCGA  
 ClusiaRosea ATGTTTACTTCCATTGTCGGTAATGTATTTGGATTCAAAGCCCTACGCGCTCTACGGCTGGAAGATTTGCGA  
 CratoxylumCochin ATGTTTACTTCCATTGTAGGTAATGTATTTGGATTCAAAGCCCTGCGTGCTCTACGGCTGGAGGATTTGCGA  
 CtenolophonEngle ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCTCTACGTCTGGAGGATTTACGA  
 DrypetesLittoral ATGTTTACTTCCATTGTGGAAATGTATTTGGGTTCAAAGCCCTACGGCTCTACGTCTAGAAGATTTGCGA  
 ElatineTriandra ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCTCTACGTCTGGAGGATTTGCGA  
 ErythroxylumNovo ATGTTGACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCTCTACGTCTGGAGGATTTGCGA  
 EuonymusHamilton ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCTCTACGTCTGGAGGATTTGCGA  
 EuphorbiaHumifus ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTGCGCGCTCTACGTCTGGAGGATTTGCGA  
 EuphroniaGuianen ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCTCTACGTCTGGAGGATTTGCGA  
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 FlueggeaVirosa ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCTCTGCGCTCTGGAGGATTTGCGA  
 GarciniaSubellip ATGTTTACTTCCATTGTCGGTAATGTATTTGGATTCAAAGCCCTACGCGCTCTACGGCTGGAGGATTTGCGA  
 GomphiaSerrata ATGTTTACTTCCATTGTGGGTAATGTATTTGGGTTCAAAGCCCTACGCGCTCTACGTCTGGAGGATTTGCGA





A. pubescens	ATCCCTCTCGCTTATACTAAAACTTTCCAAGGTCGCGCTCACGGGATTCAAGTTGAGAGAGATAAATTGAAC
A. khasayama	ATCCCTCTCGCTTATACTAAAACTTTCCAAGGTCGCGCTCACGGGATTCAAGTTGAGAGAGATAAATTGAAC
A. saxatilis	ATCCCTCTCGCTTATACTAAAACTTTCCAAGGTCGCGCTCACGGGATTCAAGTTGAGAGAGATAAATTGAAC
L. comptonii	ATCCCTGTGTGCTTATGTGAAAACTTTCCAAGGCCCGCCTCACGGGATTCAAGTTGAGAGAGATAAAGTTGAAC
H. humiriifolia	ATCCCTCTCGCTTATTCTAAAACTTTCCAAGGCCCGCCTCACGGCATTCAAGTTGAGAGAGATAAATTGAAC
I. griffithiana	-----CTTTCCAAGGTCGCGCTCACGGCATTCAAGTTGAGAGAGATAAATTGAAC
P. magnifolia	-----TAAAACTTTCCAAGGTCGCGCTCACGGCATTCAAGTTGAGAGAGATAAATTGAAC
T. sinensis	ATCCCTCTCGCTTATACTAAAACTTTCCAAGGCCCGCCTCACGGGATTCAAGTTGAGAGAGATAAATTGAAC
T. ovoidea	ATCCCTCTCGCTTATACTAAAACTTTCCAAGGCCCGCCTCACGGGATTCAAGTTGAGAGAGATAAATTGAAC
R. calophylla	ATCCCTCTCGCTTATTCTAAAACTTTCCAAGGCCCGCCTCACGGCATTCAAGTTGAGAGAGATAAATTGAAC
AcalyphaInsulana	ATCCCTATTCTTATATTCTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAT
BalanopsPancheri	ATCCCCCTGCTTATTCTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
BischofiaJavanica	ATCCCTCTCGCTTATTCTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
BruguieraGymnorh	ATTCCTCTCGCTTATTCTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
ByrsonimaCrassif	ATCCCTCTGCTTATTCTGAAAACTTTCCAAGGTCGCGCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAT
MalpighiaGlabra	ATCCCTACTGCTTATACGAAAACTTTCCAAGGCCCGCCTCATGGTATCCAAGTTGAGAGAGATAAATTGAAC
CamplostylusMann	ATTCCTCTCGCTTATTCTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
ClusiaRosea	ATCCCTCTCGCTTATACTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAAAGAGATAAATTGAAC
CratoxylumCochin	ATCCCTCTGCTTATACATAAACTTTCCAAGGTCGCGCCCATGGTATCCAAGTTGAAAGAGATAAATTGAAC
CtenolophonEngle	GTCCCTCTCGCTTATTCTAAAACTTTCCAAGGACCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
DrypetesLittoral	ATCCCTCCGGCGTATTCCAAGACTTTTCAAGGTCACCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
ElatineTriandra	ATCCCTCTGCTTATACGAAAACTTTCCAAGGACCCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
ErythroxylumNovo	ATTCCTACACTTATTCTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
EuonymusHamilton	ATCCCCCTGCTTATTCTAAAACTTTTCAAGGCCCGCCGCATGGTATCCAAGTTGAGAGAGATAAATTGAAC
EuphorbiaHumifus	ATCCCTCTGCTTATACATAAACTTTCCAAGGTCACCTCATGGTATCCAAGTTGAAAGAGATAAATTGAAC
EuphorbiaGuianen	ATCCCCCTGCTTATATTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAAAGAGATAAATTGAAC
FlourcutiaIndica	ATCCCCCTGCTTATACATAAACTTTCCAAGGCCCGCCTCACGGCATCCAAGTTGAGAGAGATAAATTGAAC
FlueggeaVirosa	ATCCCTCTGCTTATTCTAAAACTTTCCAAGGCCCGCCTCACGGCATCCAAGTTGAGAGAGATAAATTGAAC
GarciniaSubellip	ATCCCTACGGCTTATACATAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAAAGAGATAAATTGAAC
GomphiaSerrata	ATCCCTACTGCTTATATTAAAACTTTCCAAGGACCCCATGGTATCCAAGTTGAGAGAGATAAATTGAAC
HumiriaBalsamife	ATCCCTCTGCTTATTCTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
IonidiumCommune	ATCCCTCTGCTTATACTAAAACTTTTCAAGGTCGCGCTCATGGTATCCAAGTTGAGAGAGATAAATTGAAC
IrvingiaMalayana	ATCCCTACTGCTTATACATAAACTTTCCAAGGTCGCGCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
IxonanthesReticu	ATCCCTCTCGCTTATTCTAAAACTTTCCAAGGACCGCCTCATGGTATCCAAGTTGAAAGAGATAAATTGAAC
JatrophaIntegerr	ATCCCTACTGCTTATACATAAACTTTCCAAGGCCCGCCTCATGGTATCCAAGTTGAGAGAGATAAATTGAAC
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LacistemaAggrega	ATCCCTCTGCTTATTCTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
LunaniaParviflor	ATCCCTCTGCTTACTCTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAT
ManihotEsculenta	ATCCCTCTGCTTATTCTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
OxalisCorniculat	ATCCCTCTGCTTATACTAAAACTTTTCAAGGCCCGCCTCATGGTATTCAGGTTGAGAGAGATAAGTTGAAC
PassifloraQuadra	ATCCCTCTGCTTATACATAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
PhyllanthusFlexu	ATCCCTCTGCTTATTCTAAAACTTTCCAAGGCCCGCCTCACGGCATCCAAGTTGAGAGAGATAAATTGAAC
PimelodendronGri	ATCCCTACTGCTTATATTAAAACTTTCCAAGGCCCGCCTCATGGTATCCAAGTTGAGAGAGATAAATTGAAC
RicinusCommunis	GTCCCTCTGCTTATACTAAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
SacoglottisSp	ATCCCTACTTCTTATACATAAACTTTCCAAGGCCCGCCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC
ScageaOligostemo	ATCCCCCTGCTTATTCTAAAACTTTCCAAGGCCCGCCTCACGGCATCCAAGTTGAGAGAGATAAATTGAAC
Trigonibolivian	ATCCCCACTTCTTATATTAAAACTTTCCAAGGTCGCGCTCATGGCATCCAAGTTGAAAGAGATAAATTGAAC
AlbiziaTomenosaJ	ATCCCTCTCTTCTATTCTAAAACTTTCCAAGGTCGCGCTCACGGCATCCAAGTTGAGAGAGATAAATTGAAC
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QuercusRubra	ATCCCTACTTCTTATTCTAAAACTTTCCAAGGTCGCGCTCATGGCATCCAAGTTGAGAGAGATAAATTGAAC

L. suffruticosum AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
 H. clevelandii AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
 H. disjunctum AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
 L. berlandieri AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
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 H. drymarioides AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
 H. micranthum AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
 H. castaneifolia AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTACGGTAGA  
 S. digynum AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
 L. oligophyllum AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
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 L. volkensii AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTACGGTAGA  
 L. flavum AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
 L. catharticum AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
 L. arboreum AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
 R. linoides AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
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 L. comptonii AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
 H. humiriifolia AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTACGGTAGA  
 I. griffithiana AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTACGGTAGA  
 P. magnifolia AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTACGGTAGA  
 T. sinensis AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
 T. ovoidea AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCGAACTGGGCTTATCCGCTAAGAATTATGGTAGA  
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 LacistemaAggrega AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCTAAATGGGCTTATCCGCTAAGAATTACGGTAGA  
 LunaniaParviflor AAGTATGGTCGCCCCCTATTGGGCTGTACTATTAACCTAAATGGGCTTATCCGCTAAGAATTACGGTAGA





H. micranthum ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCAATTTATAAAGCACAGGCCGAAACAGGGGAAATC  
 H. castaneifolia ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCACTTTATAAAGCACAGGCCGAAACAGGTGAAATC  
 S. digynum ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCTATTTATAAATCACAGGCCGAAACAGGGGAAATC  
 L. oligophyllum ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCTATTTATAAATCACAGGCCGAAACAGGGGAAATC  
 H. neblinae ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCACTTTATAAAGCACAGGCCGAAACAGGTGAAATC  
 H. mystax ATGCGTTGGAGAGACCGTTTCTTTTNGTGCCGAAGCAATTTATAAAGCACAGGCCGAAA-----  
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 L. volkensii ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCACTTTATAAATCACAGGCCGAAACAGGGGAAATC  
 L. flavum ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCTATTTTATAAAGCACAGGCCGAAACAGGGGAAATC  
 L. catharticum ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCTATTTATAAATCACAGGCCGAAACAGGGGAAATC  
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 A. khasayama ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCAATTTATAAATCACAGGCCGAAACAGGGGAAATC  
 A. saxatilis ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCAATTTATAAATCACAGGCCGAAACAGGGGAAATC  
 L. comptonii ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCTATTTATAAATCACAGGCCGAAACAGGGGAAATC  
 H. humiriifolia ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCACTTTATAAAGCACAGGCCGAAACAGGTGAAATC  
 I. griffithiana ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCACTTTATAAAGCACAGGCCGAAACAGGTGAAATC  
 P. magnifolia ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCACTTTTATAAAGCACAGGCCGAAACAGGTGAAATC  
 T. sinensis ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCAATTTATAAATCACAGGCCGAAACAGGGGAAATC  
 T. ovoidea ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCAATTTATAAATCACAGGCCGAAACAGGGGAAATC  
 R. calophylla ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCACTTTATAAAGCACAGGCCGAAACAGGTGAAATC  
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 EuphorbiaHumifus ATGCGTTGGAGAGACCGTTTCTTATTTTGTGCCGAAGCAATTTTATAAATCACAGGCCGAAACAGGTGAAATC  
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H. obtusifolia TTGGGAGCTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
H. neocaledonic TTGGGAGCTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
H. penicillanth TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
H. spicata TTGGGAGCTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
H. planchoni TTGGGAGCTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
H. orientalis TTGGGAGCTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
H. longipes TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
L. volkensii TTGGGAGTTCCTATTGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
L. flavum TTGGGAGTTCCTATCATAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
L. catharticum TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
L. arboreum TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
R. linoides TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
R. schomburgkii TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
R. monsalveae TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
A. pubescens TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
A. khasayama TTGGGAGCTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
A. saxatilis TTGGGAGCTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
L. comptonii TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
H. humiriifolia TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
I. griffithiana TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
P. magnifolia TTGGGAACCTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
T. sinensis TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
T. ovoidea TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
R. calophylla TTGGGAGTTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
AcalyphaInsulana TTAGGAGCTCCTATCGTAATGCATGACTACTTAACAGGGGGTTCACTGCTAATACTAGCTTGGCTCATTAT  
BalanopsPancheri TTGGGAGTTCCTATCGTAATGCATGATTACTTAACAGGAGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
BischofiaJavanica TTAGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
BruguieraGymnorh TTGGGAGCTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
ByrsonimaCrassif TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
MalpighiaGlabra TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
CampostylusMann TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
ClusiaRosea TTGGGAGCTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
CratogeomysCochin TTAGGAGTTCCTATCGTAATGCATGACTACTTAACAGGAGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
CtenolophonEngle TTGGGAGTTCCTATAGTAATGCATGACTACTTAACAGGGGGTTCACTGCTAATACTAGCTTGGCTCATTAT  
DrypetesLittoral TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
ElatineTriandra TTAGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGTTCACTGCTAATACTAGCTTGGCTCATTAT  
ErythroxylumNovo TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
EuonymusHamilton CTGGGTGTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
EuphorbiaHumifus TTAGGAGTTCCTATTGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
EuphorbiaGuianen TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
FlacourtiaIndica TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
FlueggeaVirosa TTGGGAGTTCCTATCGTCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
GarciniaSubellip TTAGGAGCTCCTATCGTAATGCATGACTACTTAACAGGAGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
GomphiaSerrata TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGAGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
HumiriaBalsamife TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
IonidiumCommune TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
IrvingiaMalayana TTGGGAGTTCCTATTGTAATGCATGACTACTTAACAGGAGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
IxnanthesReticu TTAGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGTTCACTGCTAATACTAGCTTGGCTCATTAT  
JatrophaIntegerr TTAGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
KiggelariaAfrica TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
LacistemaAggrega TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
LunaniaParviflor TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
ManihotEsculenta TTAGGAGCTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
OxalisCorniculat TTGGGAGCTCCTATCGTAATGCACGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
PassifloraQuadra TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGAGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
PhyllanthusFlexu TTGGGAGTTCCTATCGTCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
PimelodendronGri TTAGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
RicinusCommunis TTAGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
SacoglottisSp TTGGGAGCTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
ScageaOligostemo TTGGGAGTTCCTATCGTAATGCATGATTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
TrigoniasBolivian TTGGGAGTTCCTATCGTAATGCATGATTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
AlbiziaTomenosaj TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
KerriaJaponica TTGGGAGTTCCTATCGTAATGCATGATTACTTAACAGGAGGATTCACTGCTAATACTAGCTTGGCTCATTAT  
DatiscaCannabina TTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGTTCACTGCTAATACTAGCTTGGCTCATTAT  
QuercusRubra CTGGGAGTTCCTATCGTAATGCATGACTACTTAACAGGGGGTTCACTGCTAATACTAGCTTGGCTCATTAT







L. pratense	GTAGGTAAACTGGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTATTGCGCGACGATTTTGTGTA
L. narbonense	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTATTGCGCGACGATTTTGTGTA
C. selaginoides	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
L. hirsutum	GTAGGTAAACTTGAAGGGGAAAGAGAAATCACTTTGGGCTTTGTTGATTTACTGCGCGACGATTTTGTGTA
L. hypericifolium	-----
L. usitatissimum	GTAGGTAAACTTGAAGGTGAAAGAGAGATCACTTTGGGCTTTGTTGATTTATTACGCGACGATTTTGTGTA
L. nervosum	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTATTGCGCGACGATTTTGTGTA
L. bienne	GTAGGTAAACTTGAAGGTGAAAGAGAGATCACTTTGGGCTTTGTTGATTTATTACGCGACGATTTTGTGTA
L. monogynum	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTATTACGCGACGATTTTGTGTA
L. suffruticosum	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
H. clevelandii	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
H. disjunctum	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
L. berlandieri	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
L. rupestre	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
D. pentagyna	GTAGGTAAACTTGAAGGGGAAAGAGACATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
R. indica	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAC
H. drymarioides	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTATTGCGTGATGATTTTATTGAA
H. micranthum	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
H. castaneifolia	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
S. distans	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
L. oligophyllum	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTATTGCGCGACGATTTTGTGTA
H. neblinae	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
H. mystax	-----
H. busseana	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
H. gabunensis	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
H. obtusifolia	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
H. neocaledonic	GTAGGTAAACTTGAAGGGGAAAGAGACATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
H. penicillanth	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
H. spicata	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
H. planchoni	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
H. orientalis	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
H. longipes	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
L. volkensii	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
L. flavum	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
L. catharticum	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
L. arboreum	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
R. linoides	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
R. schomburgkii	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTTCTGATGATTTTATTGAA
R. monsalveae	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
A. pubescens	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
A. khasayama	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
A. saxatilis	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
L. comptonii	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
H. humiriifolia	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
I. griffithiana	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
P. magnifolia	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
T. sinensis	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
T. ovoidea	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
R. calophylla	GTAGGTAAACTTGAAGGGGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
AcalyphaInsulana	GTAGGTAAACTTGAAGGGGAAAGAGAAATCACTTTGGGCTTTGTTGATTTATTGCGTGATGATTTTATTGAA
BalanopsPancheri	GTAGGTAAACTTGAAGGGGAAAGAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
BischofiaJavanica	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
BruguieraGymnorhiza	GTAGGTAAACTTGAAGGGGAAAGAGAAATCACTTTGGGCTTTGTTGATTTACTTCTGATGATTTTATTGAA
ByrsonimaCrassifolia	GTAGGTAAACTTGAAGGGGAAAGAGAGATATCACTTTAGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
MalpighiaGlabra	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTAGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
CampostylusMannii	GTAGGTAAACTTGAAGGGGAAAGAGAGATATCACTTTGGGCTTTGTTGATTTACTACGTGATGATTTTGTGTA
ClusiaRosea	GTAGGTAAACTTGAAGGTGAAAGAGAGATCACTTTGGGCTTTGTTGATTTACTGCGTTGGATTTTATTGAA
CratogeomysCochin	GTAGGTAAACTTGAAGGGGAAAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
CtenolophonEngleri	GTAGGTAAACTTGAAGGGGAAAGAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
DrypetesLittoralis	GTAGGTAAACTTGAAGGGGAAAGAGAGATATCACTTTGGGCTTTGTTGATTTACTCCGTGATGATTTTATTGAA
ElatineTriandra	GTAGGTAAACTTGAAGGGGAAAGAGAAATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
ErythroxylumNovum	GTAGGTAAACTTGAAGGTGAAAGAGAGATATCACTTTGGGCTTTGTTGATTTACTTCTGATGATTTTATTGAA
EuonymusHamiltonii	GTAGGTAAACTTGAAGGGGAAAGAGAGATATCACTTTAGGCTTTGTTGATTTACTACGTGATGATTTTGTGTA
EuphorbiaHumifusa	GTAGGTAAAGCTTGAAGGGGAAAGAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
EuphorbiaGuianensis	GTAGGTAAACTTGAAGGGGAAAGAGAGATATCACTTTAGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
FlacourtiaIndica	GTAGGTAAACTTGAAGGGGAAAGAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTGTGTA
FlueggeaVirosa	GTAGGTAAACTTGAAGGGGAAAGAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTGATGATTTTATTGAA
GarciniaSubellipica	GTAGGTAAACTTGAAGGTGAAAGAGAGATATCACTTTGGGCTTTGTTGATTTACTGCGTTGGATTTTATTGAA



R. schomburgkii	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTATCTCGCCCGTGGCTTCG
R. monsalveae	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTCTCGCCCGTGGCTTCG
A. pubescens	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTCTCGCCCGTGGCTTCG
A. khasayama	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTCTCGCCCGTGGCTTCG
A. saxatilis	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTCTCGCCCGTGGCTTCG
L. comptonii	AAAGATAGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGCGCTCTACCTGTAGCTTCG
H. humiriifolia	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTCTCGCCCGTGGCTTCG
I. griffithiana	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTCTCGCCCGTGGCTTCG
P. magnifolia	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTCTCGCCCGTGGCTTCG
T. sinensis	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTCTCGCCCGTGGCTTCG
T. ovoidea	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTCTTGCCCGTGGCTTCG
R. calophylla	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTCTCGCCCGTGGCTTCG
AcalyphaInsulana	AAAGACCGAAGCCGTGGCATTATTTATTCAGCAAGATTGGGTCTCTCTACCAGGTGTCTCGCTGTAGCTTCA
BalanopsPancheri	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCGGAGTTCTGCCCGTGGCTTCA
BischofiaJavanica	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTTTACCAGGTGTTATTTCCCGTGGCTTCA
BruguieraGymnorh	AAAGATCGAGGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTATACCCGTGGCTTCA
ByrsonimaCrassif	AAAGATAGAAGCCGCGGATTTATTTTCACTCAAGATTGGGTCTCTCTACCTGGTGTATATACCCGTGGCTTCA
MalpighiaLabra	AAAGATAGAAGCCGCGGATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTATACCTGTGGCTTCG
CamplostylusMann	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTATACCCGTGGCTTCA
ClusiaRosea	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCCGGTGTCTTCTCTGTGGCTTCA
CratoxylumCochin	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTATACCCGGTGTCTTGCCGTGGCTTCC
CtenolophonEngle	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGCGTCTTGCCCGTGGCTTCA
DrypetesLittoral	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTTTGCCCGTGGCTTCT
ElatineTriandra	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCTGGGTGTTCTACCTGTGGCTTCT
ErythroxylumNovo	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTCTGCCGTGGCTTCA
EuonymusHamilton	AAAGACCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTCTGCTGTGGCTTCT
EuphorbiaHumifus	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTTGTGCTGGTGTCTTACCTGTAGCTTCA
EuphorbiaGuianensis	AAAGATCGAAGTCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCGGGTCTTGCCCGTGGCTTCC
FlacourtiaIndica	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTATTTCCCGTGGCTTCG
FlueggeaVirens	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTATGCCGGGTGTTATACCTGTGGCTTCA
GarciniaSubellip	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTCTTCTGTAGCTTCA
GomphiaSerrata	AAAGATCGAAGTCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTATCCCGTGGCTTCA
HumiriaBalsamifera	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTATACCCGTGGCTTCA
IonidiumCommune	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAGATTGGGTCTCTCTACCCGGTGTCTTGCCGTGCTTCCG
IringiaMalayana	AAAGATCGAAGCCGTGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCTGGTGTGTTTGCCCGTGGCTTCC
IxonanthesReticulata	AAAGATCGAAGCCGCGCATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTATACCCGTGGCTTCC
JatrophaIntegrifolia	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTGTTGCTGTAGCTTCA
KigeliaAfrica	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTTTACCAGGTGTTCTGCGTGTGGCTTCG
LacistemaAggregata	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTCTGCTGTGGCTTCA
LunaniaParviflora	AAAGACCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTCTGCCGTAGCTTCA
ManihotEsculenta	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTCTGCTGTAGCTTCA
OxalisCorniculata	AAAGACCGAAGCCGTGGGATTTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTCTGCCGTGGCTTCA
PassifloraQuadrata	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTATTTCCCGTGGCTTCA
PhyllanthusFlexuosus	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTATGCCGGTGTGTTTACCTGTGGCTTCA
PimelodendronGri	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCTGGTGTGTTTACCTGTAGCTTCA
RicinusCommunis	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCTGGTGTGTTCTGCTGTAGCTTCA
SacoglottisSp	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTATACCCGTGGCTTCA
ScageaOligostemo	AAAGATCGAAGTCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCTGGTGTGTTCTGCCGTGGCTTCA
TrigonostylisBolivia	AAAGACCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCGGGTGTATACCTGTGGCTTCA
AlbiziaTomenosaj	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAGGATTGGGTCTCTCTACCAGGTGTTCTGCCGTGTGCTTCG
KerriaJaponica	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTGCCAGGTGTTTGGCTGTGGCTTCA
DatiscaCannabina	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTCTACCAGGTGTTCTGCTGTGGCTTCA
QuercusRubra	AAAGATCGAAGCCGCGGTATTTATTTTCACTCAAGATTGGGTCTCTTTACCAGGTGTTCTGCCCGTGGCTTCC

L. lewisii GGGGGTATTACAGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCGGTATTACAATTTGGC  
L. perenne GGGGGTATTACAGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCGGTATTACAATTTGGC  
L. pratense GGGGGTATTACAGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCGGTATTACAATTTGGC  
L. narbonense GGGGGTATTACAGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCGGTATTACAATTTGGC  
C. selaginoides GGGGGTATTACAGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCGTATTACAATTTGGC  
L. hirsutum GGGGGTATTACATGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCGTATTACAATTTGGA  
L. hypericifolium-----

L. usitatissimum GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 L. nervosum GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 L. bienne GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 L. monogynum GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 L. suffruticosum GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. clevelandii GGCGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCTGTATTACAATTTGGC  
 H. disjunctum GGCGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCTGTATTACAATTTGGC  
 L. berlandieri GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCTGTATTACAATTTGGC  
 L. rupestre GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCTGTATTACAATTTGGC  
 D. pentagyna GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 R. indica GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. drymarioides GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCTGTATTACAATTTGGC  
 H. micranthum GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCTGTATTACAATTTGGC  
 H. castaneifolia GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 S. digynum GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCTGTATTACAATTTGGC  
 L. oligophyllum GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. neblinae GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. mystax -----  
 H. busseana GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. gabunensis GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. obtusifolia GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. neocaledonic GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. penicillanth GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. spicata GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. planchoni GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. orientalis GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 H. longipes GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 L. volkensii GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 L. flavum GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 L. catharticum GGGGGTATTACGTTTGGCATATGCCCGCTTTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 L. arboreum GGGGGTATTACGTTTGGCATATGCCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 R. linoides GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 R. schomburgkii GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 R. monsalveae GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 A. pubescens GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 A. khasayama GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 A. saxatilis GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 L. comptonii GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCTGTATTACAATTTGGC  
 H. humiriifolia GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 I. griffithiana GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 P. magnifolia GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 T. sinensis GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 T. ovoidea GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 R. calophylla GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 AcalyphaInsulana GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 BalanopsPancheri GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 BischofiaJavanic GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 BruguieraGymnorh GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 ByrsonimaCrassif GGAGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 MalpighiaGlabra GGAGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 CamptostylusMann GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 ClusiaRosea GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCTGTATTACAATTTGGT  
 CratogeomysCochin GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 CtenolophonEngle GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 DrypetesLittoral GGGGGTATTACGTTTGGCATATGCCCGCTCTGACAGAGATCTTTGGAGATGATTCTGTATTACAATTTGGT  
 ElatineTriandra GGAGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 ErythroxylumNovo GGAGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 EuonymusHamilton GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 EuphorbiaHumifus GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 EuphoniaGuianen GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 FlacourtiaIndica GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 FlueggeaVirosa GGAGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 GarciniaSubellip GGAGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 GomphiaSerrata GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 HumiriaBalsamife GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGT  
 IonidiumCommune GGAGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC  
 IrvingiaMalayana GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCTGTATTACAATTTGGT  
 IxonanthesReticu GGGGGTATTACGTTTGGCATATGCCTGCTCTGACAGAGATCTTTGGAGATGATTCCGTATTACAATTTGGC



[illegible]

L. comptonii	GGTGGAACTTTAGGTCACCCCTTGGGGAAATGCACCTGGTGCTGTAGCTAATCGAGTAGCTCTAGAAGCATGT
H. humiriifolia	GGAGGAACTTTAGGTACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
I. griffithiana	GGAGGAACTTTAGGTACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
P. magnifolia	GGAGGAACTTTAGGTACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
T. sinensis	GGTGGAACTTTAGGTCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGGGTAGCTCTAGAAGCATGT
T. ovoidea	GGTGGAACTTTAGGTACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGGGTAGCTCTAGAAGCATGT
R. calophylla	GGAGGAACTTTAGGTACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
AcalyphaInsulana	GGAGGAACTTTAGGGCATCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
BalanopsisPancheri	GGAGGAACTTTAGGGCACCCGTGGGGAAATGCACCTGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
BischofiaJavanic	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
BruguieraGymnorh	GGAGGAACTTTAGGACACCCCTTGGGGAAATGCACCCAGGTGCTGTAGCTAATCGAGTAGCTCTAGAAGCATGT
ByrsonimaCrassif	GGAGGAACTTTGGGGCACCCCTTGGGGAAATGCACCTGGTGCCGTAGCTAATCGAGTAGCTCTCGAAGCATGT
MalpighiaGlabra	GGAGGAACTTTGGGGCACCCCTTGGGGAAATGCACCCAGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
CamptostylusMann	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
ClusiaRosea	GGAGGAACTTTAGGGCATCCTTGGGGAAATGCACCTGGTGCTGTAGCTAATCGAGTAGCTCTAGAAGCATGT
CratoxylumCochin	GGAGGAACTTTAGGGCATCCTTGGGGAAATGCACCTGGTGCTGTAGCTAATCGAGTTGCTCTAGAAGCATGT
CtenolophonEngle	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCTGTAGCTAATCGAGTAGCTCTAGAAGCATGT
DrypetesLittoral	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
ElatineTriandra	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCTGGTGCTGTAGCTAATCGAGTCGCTCTAGAAGCATGT
ErythroxylumNovo	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
EuonymusHamilton	GGAGGAACTTTAGGGCATCCTTGGGGAAACGACCCCTGGTGCTGTAGCTAATCGAGTAGCTCTAGAAGCATGT
EuphorbiaHumifus	GGCGGAACTTTAGGGCACCCCTTGGGGAAATGCACCTGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
EuphroniaGuianen	GGAGGAACTTTAGGACACCCGTGGGGCAATGCACCAGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
FlacourtiaIndica	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTGCTAATCGAGTAGCTCTAGAAGCATGT
FlueggeaVirosa	GGTGGAACTTTAGGGCACCCCTTGGGGAAATGCACCTGGTGCCGTAGCTAACCAGTAGCTCTAGAAGCATGT
GarciniaSubellip	GGAGGAACTTTAGGCCATCCTTGGGGAAATGCACCTGGTGCTGTAGCTAATCGAGTAGCTCTAGAAGCATGT
GomphiaSerrata	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
HumiriaBalsamife	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
IonidiumCommune	GGAGGAACTTTGGGGCACCCCTTGGGGAAATGCACCCAGGTGCCGTGCTAATCGAGTAGCTCTAGAAGCATGT
IrvingiaMalayana	GGAGGAACTTTGGGGCATCCTTGGGGAAACGACCCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
IxonanthesReticu	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCGCCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
JatrophaInteger	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCGCCGTGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
KiggelariaAfrica	GGAGGGACTTTAGGTACCCCTTGGGGAAATGCACCCGGTGCCGTGCTAATCGAGTAGCTCTCGAAGCATGT
LacistemaAggrega	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTGCTAATCGAGTAGCTCTAGAAGCATGT
LunaniaParviflor	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTGCTAATCGAGTAGCTCTGGAAGCATGT
ManihotEsculent	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
OxalisCorniculat	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCTGGTGCTGTAGCTAATCGAGTGGCTTTAGAAGCATGT
PassifloraQuadra	GGAGGCACTTTAGGGCACCCCTTGGGGAAATGCACCTGGTGCCGTGCTAATCGAGTAGCTCTAGAAGCATGT
PhyllanthusFlexu	GGTGGAACTATAGGGCACCCCTTGGGGAAATGCACCCAGGTGCTGTAGCTAACCAGTAGCTCTAGAAGCATGT
PimelodendronGri	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCTGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
RicinusCommunis	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
SacoglottisSp	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
ScageaOligostemo	GGAGGAACTTTAGGGCACCCGTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCCTTGT
TrigonialBolivian	GGGGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
AlbiziaTomenosaJ	GGCGGAACTTTAGGACACCCCTTGGGNAATGCACCCGGTGNNGTAGCTAATCGAGTAGCTCTGGAAGCATGT
KerriaJaponica	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
DatiscaCannabina	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCCGTAGCTAATCGAGTAGCTCTAGAAGCATGT
QuercusRubra	GGAGGAACTTTAGGGCACCCCTTGGGGAAATGCACCCGGTGCTGTAGCTAATCGAGTAGCTCTAGAAGCATGT

H. disjunctum GTACAAGCGCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCG-----TAT  
L. berlandieri GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
L. rupestre GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
D. pentagyna GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
R. indica GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
H. drymarioides GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
H. micranthum GTACAAGCGCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
H. castaneifolia GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
S. digynum GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
L. oligophyllum GTACAAGCTCGKAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
H. neblinae GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
H. mystax -----TAT  
H. busseana GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
H. gabunensis GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
H. obtusifolia GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
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H. penicillanth GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
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H. planchoni GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
H. orientalis GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
H. longipes GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
L. volkensii -----TAT  
L. flavum GTACAAGCTCGGAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
L. catharticum GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
L. arboreum GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAA---  
R. linoides GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
R. schomburgkii GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGCGGCTAGCAAATAT  
R. monsalveae GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAA---  
A. pubescens GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
A. khasayama GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
A. saxatilis GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCG-----TAT  
L. comptonii GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
H. humiriifolia GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTC-----  
I. griffithiana GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAAT-----TAT  
P. magnifolia GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
T. sinensis GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAAATAT  
T. ovoidea GTACAAGCTCGTAATGAGGGACGTG-----TAT  
R. calophylla GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGCGGCTAGCAA---  
AcalyphaInsulana GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTAGGCTAGTAAA---  
BalanopsPancheri GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGTAAA---  
BischofiaJavanic GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
BruguieraGymnorh GTACAAGCTCGGAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGGGCTAGCAA---  
ByrsonimaCrassif GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
MalpighiaGlabra GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
CamptostylusMann GTAAAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGAGTAATGAAATTATCCGTGAGGCTAGCAA---  
ClusiaRosea GTGCAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGTAAA---  
CratogeomysCochin GTCAAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAA---  
CtenolophonEngle GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAA---  
DrypetesLittoral GTACAAGCTCGGAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
ElatineTriandra GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGTAAA---  
ErythroxylumNovo GTACAAGCTCGTAACGAAGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
EuonymusHamilton GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
EuphorbiaHumifus GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGGCAA---  
EuphoniaGuianen GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGTAAA---  
FlacourtiaIndica GTAAAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
FlueggeaVirosa GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
GarciniaSubellip GTGCAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAA---  
GomphiaSerrata GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATACGCGAGGCTAGCAA---  
HumiriaBalsamife GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
IonidiumCommune GTACAAGCTCGTAATGAAGGCGCGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
IrvingiaMalayana GTACAAGCCCCGTAAATGAAGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
IxnanthesReticu GTACAAGCTCGCAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
JatrophaIntegerr GTACAAGCTCGTAATGAAGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATTCGTGAGGCTAGCAA---  
KiggelariaAfrica GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
LacistemaAggrega GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAACGAAATTATACGTGAGGCTAGCAA---  
LunaniaParviflor GTAAAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAGGCTAGCAA---  
ManihotEsculenta GTACAAGCTCGTAATGAGGGACGTGATCTTGCTCGTGAGGGTAATGATATTATCCGTGAGGCTAGCAA---  
OxalisCorniculat GTACAAGCTCGTAATGAAGGCGGTGATCTTGCTCGTGAGGGTAATGAAATTATCCGTGAAGCTAGCAA---





H. drymarioides TATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTT-----TCAGAAAAATTTAGGTTGT  
H. micranthum TATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTT-----TCAGAAAAATGTTAGGTTGT  
H. castaneifolia TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCAATTTTG-----TTGGAAAAATGTTAGGTTAT  
S. digynum TATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTT-----TCAGAAAAATTTAGGTTGT  
L. oligophyllum TATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTT-----TCAGAAAAATGTTAGGTTGT  
H. neblinae TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCCATTTTG-----TTGGAAAAATGTTAGGTTAT  
H. mystax TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCAATTTTG-----TTGGAAAAATGTTAGGTTAT  
H. busseana TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCAATTTTG-----TTGGAAAAATGTTAGGTTAT  
H. gabunensis TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCAATTTTG-----TTGGAAAAATGTTAGGTTAT  
H. obtusifolia TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCAATTTTG-----TTGGAAAAATGTTAGGTTAT  
H. neocaledonic TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCAATTTTG-----TTGGAAAAATGTTAGGTTAT  
H. penicillanth TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCCATTTTG-----TTGGAAAAATGTTAGGTTAT  
H. spicata TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCAATTTTG-----TTGGAAAAATGTTAGGTTAT  
H. planchoni TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCAATTTTG-----TTGGAAAAATGTTAGGTTAT  
H. orientalis TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCAATTTTG-----TTGGAAAAATGTTAGGTTAT  
H. longipes TATACAGTTACTTATAATCGTGGTTTAAATGGATATGGATCAATTTTG-----TTGGAAAAATGTTAGGTTAT  
L. volkensii TATATAGTTACTTACGCTCGTGGTTTAAACGGATATGGATCCATTTT-----TCAGAAAAATGTTAGGTTGT  
L. flavum TATATAGTTACTTATGCTCGTGGTTTAAATCGATATGGATCCATTTT-----TCGGAAACATGTTGGGTTGT  
L. catharticum TATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGATCCCTTTT-----TCGGAAAAATGTTAGGTTGT  
L. arboreum -----  
R. linoides TATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGATCCATT-----AAGGTAGGTTAT  
R. schomburgkii TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCCATTTTG-----TTGGAAAAATGTTAGGTTAT  
R. monsalveae -----  
A. pubescens TATATAGTTACTTATGCTCGTGGTTTAAATGGCTATGGATCCATTTT-----TCGGAAAAATGTTAGGTTAT  
A. khasayama TATATAGTTACTTATGCTCGTGGTTTAAATGGCTATGGATCCATTTT-----TCGGAAAAATGTTAGGTTAT  
A. saxatilis TATATAGTTACTTATGCTCGTGGTTTAAATGGCTATGGATCCATTTT-----TCGGAAAAATGTTAGGTTAT  
L. comptonii TATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTT-----TCAGAAAAATGTTAGGTTGT  
H. humiriifolia TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCCATTTTG-----TTGGAAAAATGTTAGGTTAT  
I. griffithiana TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCAATTTTG-----TTGGAAAAATGTTAGGTTAT  
P. magnifolia TATACAGTTACTTATGATCGTGGTTTAAATGGATATGGATCCATTTTG-----TTGGAAAAATGTTAGGTTAT  
T. sinensis TATATACTTACTTATGCTCGTGGTTTAAATGGATATGTATCCATTTT-----TCGGAAAAATGTTGGGTTAT  
T. ovoidea TATATACTTACTTATGCTCGTGGTTTAAATGCATATGGATCCATTTT-----TCGGAAAAATGTTGGGTTAT  
R. calophylla -----  
AcalyphaInsulana TATACATTTGCTCATGATCATAGTTTAAAT-----AGATCTAGTTTG-----TTGGAAAAATGTTAGGTTAT  
BalanopsPancheri TATACATTTGCTTATGATCATGTTTAAAT-----AGATCTATTTTG-----TTGGAAAAAGTTAGGTTAT  
BischofiaJavanica TATAGACTTGTTCATGATCATGTTTAAAT-----AGATCTATTTTG-----TTGGAAAAATGTTGGGTTAT  
BruguieraGymnorh TATACACTTTCTCATGATCATGTTTAAAT-----AAATCTATTTTG-----TTGGAAAAATGAAAGGTCAT  
ByrsonimaCrassif TATACACTTGCTCATGATCATGTTTAAAT-----AGATCGATTTTG-----TTGGAAAAATGTTGGGTTAT  
MalpighiaGlabra TATATACTTGCTCAT-----GTAAAT-----AGATCTATTTTA-----TTGGAAAAATGTTAGGTTAT  
CampostylusMann TATACACTTGCTCATGATCATGTTTAAAT-----AGATCTATTTTG-----TTGGAAAAATGTTGGGTTAT  
ClusiaRosea -----TTGCTCATGGTCATGTTTAAATA-----AATAGATCCATTTTA-----TTGGAAAAATGTTAGGTTAT  
CratoxylumCochin TATATACTTGCTCATGATCATAGTTTACAT-----AGATCTGCTTTG-----TTGGAAAAATTTAAGTTAT  
CtenolophonEngle -----  
DrypetesLittoral TATACACTTGCTCATGATCATAGTTTAAAT-----AGATCTCTTTTG-----TTGGAAAAATGTTAGGTTAT  
ElatineTriandra TATACACTTGCTCATGATCATGTTTAAAT-----AGATCGATTTTG-----TTGGAAAAATGCGGGTTAT  
ErythroxylumNovo TATATACTTGTTTCATGATCATCGGTTTAAAT-----AGATCCATTTTG-----TTGGAAAAATGGAGATTAT  
EuonymusHamilton -----TTCCCATGATTATGTTTAAAT-----AGATCTATTTTG-----TTGCAAAATGTCAGTTAT  
EuphorbiaHumifus TATACATTTTCTCATGATCATAGTTTAAAT-----AGATCTATTTTG-----TTGCAAAATGAAAGGTTAT  
EuphroniaGuianen TATACGCTTGCTCATGATCATCGTTTAAAGTTAAACCGATCTATTTTGTTTAAATTTAAATATGCAAGGTTAT  
FlacourtiaIndica TATACATTTGCTCATGATCATGCTGACTTAAAT-----AGATCTATTTTG-----TTAGAAAAATGTTAGGTTAT  
FlueggeaVirosa TATAGGCTTGCTCATGAGCATGGTTTAAAT-----AAATCTATTTTG-----TTGGAAAAATGTTGGGTTAT  
GarciniaSubellip TATACACTTGCTCATGATCATGTTTAAATAGAGATAGATATATTTTA-----TTGGAAACTGTTAGGTTAT  
GomphiaSerrata TATACACTTGCTTATGATCATGTTTAAAT-----AGATCTATTTTG-----TTGGAAAAATGTTAGGTTAT  
HumiriaBalsamife TATACACTTGCTCATGATCATGTTTAAAT-----GGATCTATTTCTG-----TTGGAAAAATGTTGGGTTAT  
IonidiumCommune TATACATTTGCTCATGATCATGTTTAAAT-----AAATCTCTTTTG-----TTGGAAAAATGTTGGGTTAT  
IrvingiaMalayana TATACATTTGCTCATGATCATGTTTAAAT-----AGATCCATTTTG-----TTGGAAAGATGTTAGGTTAT  
IxonanthesReticu TATACACTTGCTCATGATCATGTTTAAAGC-----GGATCCATTTTC-----TTAGAAAAATGTTAGGTTAT  
JatrophaIntegerr TATATATTTGCTCATGATCATAGTTTAAAT-----AGATCGATTTTG-----TTGGAAAAATGTTAGGTTAT  
KiggelariaAfrica TATATATTTGCTCATGATCATGTTTAAAC-----GGATCTATTTTG-----TTGGAAAAATTTAGATTAT  
LacistemaAggrega TATACACTTGCTCATATATCATGTTTAAAT-----AGATCTATTTTG-----TTGGAAAAATGTTAGGGTAT  
LunaniaParviflor TATACGTTTGCTCATGATCGTGATTTAAAT-----AGATCGATTTTG-----TTGGAAAAATGTTAGGTTAT  
ManihotEsculenta TATACATTCGCTCATGATCATAGTTTAAAT-----AGATCTATTTTG-----TTGGAAAAATGTTAGGTTAT  
OxalisCorniculat --TGCCCTTGCTCAGCATCATTTTTTAAATAAAAAATCGATCCATTTTG-----TTGGAAAAATGTCAGATTAT  
PassifloraQuadra TATACTCTTGTTTCATGATCGTGGTTTCAAT-----AGATCGATTGTA-----TTGCAAAATATGGCTTAT  
PhyllanthusFlexu TATAGACTTGCTCATGAGCATGTTTAAAT-----AAATCTATTTTG-----TTGGAAAAATGTTAGGTTAT  
PimelodendronGri -----TCATGATCATAGTTTAAAT-----AGATCGATTTTG-----TTGGAAAAATGTTAGGTTAT  
RicinusCommunis TATACATTTGCTCATGATCATAGTTTAAAT-----AGATCTACTTTG-----TTGGAAAAATTTAGGTTAT  
SacoglottisSp TATACACTTGCTCATGATCATGTTTAAAT-----GGATCTATTTCTG-----TTGGAAAAATGTTAGGTTAT







S. digynum	TTC-----ATTATTTCTGGCTACT-----GATTCTGAACAACATATACTTTTT
L. oligophyllum	TTC-----ATTATTTCTGGCTACT-----GATTCTGAACAACATATACTTTTT
H. neblinae	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
H. mystax	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
H. busseana	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
H. gabunensis	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
H. obtusifolia	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
H. neocaledonic	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
H. penicillanth	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
H. spicata	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
H. planchoni	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
H. orientalis	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
H. longipes	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
L. volkensii	TTC-----ATTATTTCTGGCTACT-----GATTCTGAACAACATATACTTTTT
L. flavum	TTC-----ATTATTTCTGGCTACT-----GATTCTGAACAACATATACTTTTT
L. catharticum	TTC-----ATTGTTTCGCATACT-----GATTCTGAACAACATATACTTTTT
L. arboreum	-----
R. linoides	TTC-----ATTATTTCTGGCTACT-----GATTCTGAACAACATACACTTTTGT
R. schomburgkii	TTC-----ATTATTTCTGGCTACT-----GATTCTGAAAAAATACGCTTTTT
R. monsalveae	-----
A. pubescens	TTC-----ATTATTTCTGGCTACT-----GATTCTCAGCAACATACACTTTTT
A. khasayama	TTC-----ATTATTTCTGGCTACT-----GATTCTCAGCAACATACACTTTTT
A. saxatilis	TTC-----ATTATTTCTGGCTACT-----GATTCTCAGCAACATACACTTTTT
L. comptonii	TTC-----ATTATTTCTGGCTACT-----GATTCTGAACAACATATACTTTTT
H. humiriifolia	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
I. griffithiana	TTC-----ATTATTTCTGCTACT-----TATTCTGAAAAAATACACTTTTT
P. magnifolia	TTC-----ATTATTTCTGCTACT-----GATTCTGAAAAAATACACTTTTT
T. sinensis	TTC-----ATTATTTCTGGCTATT-----GATTCTGAACAACATATACTTTTT
T. ovoidea	TTC-----ATTATTTCTGGCTATT-----GATTCTGAACAACATACACTTTTT
R. calophylla	-----
AcalyphaInsulana	TTG-----ATTTTTTCTTCTAAT-----GATTCTAATCAAAATAAATTTTTT
BalanopsPancheri	TTG-----ATTATTTCTATTAAT-----GGTTCTAATCAAAATCGATTTAT
BischofiaJavanica	TTG-----ATTATTTCTGCTAAT-----AATTCTAACC AAAATCCGTTTTTG
BruguieraGymnorh	TTA-----ATTATTTCTGCAAAAT-----GATTCTAACC AAAATCCATTTCTT
ByrsonimaCrassif	TTG-----ATTATTTCTGCTAAT-----GATTCTAACC AAAATCGATTTTTT
MalpighiaGlabra	TTG-----GTTATTCCTGTAAAT-----GATTCTAACC AAAATCGATTTTTT
CamptostylusMann	TTG-----GTTATTCCTGTAAAT-----GATTCTAACC AAAATCGATTTTTT
ClusiaRosea	TTG-----ATTATTTCTGGCTAAT-----GAGTCTAACC AAAATCTATTTTTT
CratogeomysCochin	TTG-----ATTATTTCTGATAAT-----GATTATAACAAAATCTATTTTTT
CtenolophonEngle	-----
DrypetesLittoral	TTG-----ATTATTTCTGCTGAT-----GATTCTAACC AAAATCCATTTTTT
ElatineTriandra	TTG-----ATTATTTCTGCTAAT-----GATTCTAACC AAAATCCGTTTTT
ErythroxylumNovo	TTT-----CTTTTTTCTGGAAAG-----AATTCTAACC AAAATCAATTTCTT
EuonymusHamilton	TTG-----ATTATTTCTACTAAT-----TATTCTAACC AAAATCTATTTTTT
EuphorbiaHumifus	TTG-----ATTATTTCTGCTAAT-----GATTCTAACC AAAATCCATTTTTT
EuphorbiaGuianen	TTG-----ATTAGTTCTGTAAAT-----GATTCTAATCAAAATCCATTTTTT
FlacourtiaIndica	TTG-----ATTATTTCTGCTAAT-----GATTCTAACC AAAATCCATTTTTT
FlueggeaVirosa	TTG-----ATTATTTCTGGCTAAT-----AATTCTAACC AAAACTCGTTTTT
GarciniaSubellip	TTG-----ATTATTTCTGCTAAT-----GAGTCTAACC AAAATAGATTTTTT
GomphiaSerrata	TTG-----ATTATTTTCTGCTAAT-----GGTCTAACC AAAATCGATTTAT
HumiriaBalsamife	TTG-----ATTATTTCTGCTAAT-----GATTCTAACC AAAATCCATTTTTT
IonidiumCommune	TTG-----ATTTTTTCTATTAAT-----GATTCTAATCAAAATCCGTTTTT
IrvingiaMalayana	TTG-----ATTGTTTCTACTAAT-----AATTCTAACC AAAATTCATTTTTT
IxonanthesReticu	TTA-----ATTATTTTTTCTAAT-----GATTTTAACC AAAATACATTTTTT
JatrophaIntegerr	TTGGTTTTTTTTTCTTTTTTCTGCTAAT-----GATTCTAACC AAAATTCATTTTTT
KiggelariaAfrica	TTG-----ATTAGTTCTATTAAT-----GATTCTAATCAAAATAAATTTTTT
LacistemaAggrega	TTG-----ATTATTTCTGCTAATAATTCTAACC AAAATAGAAATTCATTTTTT
LunaniaParviflor	TTA-----ATTATTTCTGCTAAT-----GATTCTAACC AAAATCGATTTTTT
ManihotEsculenta	TTG-----ATTCTTTCTGCTAAT-----GATTCTAACC AAAATCTATTTTTT
OxalisCorniculat	TTG-----ATTTTTTCTACTAAT-----GATTCTGCTAACC AAAATCCGTTTTT
PassifloraQuadra	TTT-----ATTTTGCTTTTAAAT-----GATTCTAACC AAAATCGATTTTTT
PhyllanthusFlexu	TTG-----ATTATTTCTGCTAAT-----AATTCTAACC AAAATCGATTTTTT
PimelodendronGri	TTG-----ATTATTTCTGCTAAT-----GATTCTAACC AAAATCAATTTTTT
RicinusCommunis	TTG-----ATTATTTCTGCTAAT-----GATTCTAATCAAAATCCATTTTTT
SacoglottisSp	TTG-----ATTATTTCTGCTAAT-----GATTCTAACC AAAATCCATTTTTT
ScageaOligostemo	TTG-----ATTTTTTTTACTAAA-----GATTCTAACC AAAATTCGTTTTT
TrigoniasBolivian	TTG-----ATTAGTTCTATTAAT-----GATTCTAATCAAAATAAATTTTTT
AlbiziaTomenosaj	TTG-----ATTATTTCTGCTAAT-----AATTCTAACC AAAATCCATTTTTG





H. mystax CTACGATTAGTATCCTCT-----TTAGAAAGGTCAAAG-----ATAGTCAAATCCCATAATTTACGATCA  
H. busseana CTACGATTAGTATCCTCT-----TTAGAAAGGTCAAAG-----ATAGTCAAATCCCATAATTTACGATCA  
H. gabunensis CTACGATTAGTATCCTCT-----TTAGAAAGGTCAAAG-----ATAGTCAAATCCCATAATTTACGATCA  
H. obtusifolia CTACGATTAGTATCCTCTTTAGATTAGAAAGGTCAAAG-----ATAGTCAAATCCCATAATTTACGATCA  
H. neocaledonic CGACGATTAGTATCCTCT-----TTAGAAAGGTCAGAG-----ATAGTCAAATCCCATAATTTACGATCA  
H. penicillanth CGACGATTAGTATCCTCT-----TTAGAAAGGTCAGAG-----ATAGTCAAATCCCATAATTTACGATCA  
H. spicata CTACGATTAGTATCCTCTTTAGATTAGAAAGGTCAAAG-----ATAGTCAAATCCCATAATTTACGATCA  
H. planchoni CTACGATTAGTATCCTCTTTAGATTAGAAAGGTCAAAG-----ATAGTCAAATCCCATAATTTACGATCA  
H. orientalis CTACGATTAGTATCCTCT-----TTAGAAAGGTCAAAG-----ATAGTCAAATCCCATAATTTACGATCA  
H. longipes CTACGATTAGTATCCTCT-----TTAGAAAGGTCAAAG-----ATAGTCAAATCCCATAATTTACGATCA  
L. volkensii CTACGATTAGTATCCGCT-----TTAGAAAGGTTAGAA-----GTAGTCAAATCCCAAAATTTACGATCA  
L. flavum CTCCGATTAGTATCCGCT-----TTAGAAAGGTTAGAA-----GTAGTCAAATCCCAAAATTTACAATCA  
L. catharticum TTACGATTAGTATCCGCT-----TTAGAAAGCTTAGAA-----GTAGTCAAATCCCAAAATTTACGATCA  
L. arboreum -----  
R. linoides CTACGATTAGTATCCACT-----TTAGAAAGCTTCGAA-----GTAGTCAAATCCCAAAATTTACGATCA  
R. schomburgkii CTACGATTAGTATCCTCT-----TTAGAAAGGTCAGAG-----ATAGTAAAATCCCATAATTTACGATCA  
R. monsalveae -----  
A. pubescens CTACGATTAGTATCCTCT-----TTAGAAAGGTTAGAA-----ATAGTCAAATCCCAAAATTTACGATCA  
A. khasayama CTACGATTAGTATCCTCT-----TTAGAAAGGTTAGAA-----ATAGTCAAATCCCAAAATTTACGATCA  
A. saxatilis CTACGATTAGTATCCTCT-----TTAGAAAGGTTAGAA-----ATAGTCAAATCCCAAAATTTACGATCA  
L. comptonii CTACGATTAGTATCCGCT-----TTAGAAAGGTTAGAA-----GTAGTCAAATCCCAAAATTTACGATCA  
H. humiriifolia CTACGATTAGTATCCTCT-----TTAGAAAGGTCAGAG-----ATAGTCAAATCCCATAATTTACGATCA  
I. griffithiana CTACGATTAGTATCCTCT-----TTAGAAAGGTCAGAG-----ATAGTAAAATCCCATAATTTACGATCA  
P. magnifolia CTACGATTAGTATCCTCT-----TTAGAAAGGTCAGAG-----ATAGTGAATCCCAATTTACGATCA  
T. sinensis CTACGATTAGTATCCTCTTTAGATTAGCCAGCTTAGAA-----ATAGTCAAATCCCAAAATTTACGATCA  
T. ovoidea CTACGATTAGTATCCTCTTTAGATTAGACAGCTTAGAA-----ATAGTCAAATCCCAAAATTTACGATCA  
R. calophylla -----  
AcalyphaInsulana CTACGATTAGTATCCTCT-----TTAGAAAGGTCAGAG-----ATAGTCAAATCTCATAAATTTACGATCA  
BalanopsPancheri CTACGATTGTATTCTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
BischofiaJavanic CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
BruguieraGymnorh CTACAATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTCAAATCTCATAATTTACGATCA  
ByrsonimaCrassif CCTCGATTAGTATCGTCC-----TTAGAAAGGTCAGGA-----ATACAAAAATCCCTAATTTACGATCA  
MalpighiaGlabra CCTCGATTAAATATCGTCC-----TTAGAAAGGTCAGGA-----ATACAAAAATCCCTAATTTACGATCA  
CamptostylusMann CCTCGATTCTGATCGTCC-----TTAGAAAGGTCAGGA-----ATACAAAAATCCCTAATTTACGATCA  
ClusiaRosea CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTCAAATTTTACGATCA  
CratogeomysCochin CTACGATTATGCTTCT-----TTAGAAAGGTCAGAA-----ATAGTCAAATTTTACGATCA  
CtenolophonEngle -----  
DrypetesLittoral CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
ElatineTriandra CTGCGATTATATCTGCTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
ErythroxylumNovo CTCCAATTAGTATCTTCT-----TTAGACGGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
EuonymusHamilton CTCCGATTGACCTTCT-----TTAGAAAGGTCAGAA-----ATAGTCAAATCTCATAATTTACGATCA  
EuphorbiaHumifus CGACAATTAGGATCTTCT-----TTAGAAAGGTCAGAG-----ATAGTAAAATCGCATAAATTTACGATCA  
EuphorbiaGuianen TCCCGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGGGAATCTCATAATTTACGATCA  
FlacourtiaIndica TTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
FlueggeaVirosa CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTCAAATCTCATAATTTACGATCA  
GarciniaSubellip CTACGATTGAACTTCT-----TTAGAAAGGTCAGAA-----ATAGTCAAATCTCATAATTTACGATCA  
GomphiaSerrata CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
HumiriaBalsamife CTACGATTAGTATTTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
IonidiumCommune CGACGATTAGTATCCTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
IrvingiaMalayana TTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
IxnanthesReticu TCACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
JatrophaIntegerr CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAG-----ATAGTAAAATCTCATAATTTACGATCA  
KiggelariaAfrica CGACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
LacistemaAggrega CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
LunaniaParviflor CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
ManihotEsculenta CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
OxalisCorniculat CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
PassifloraQuadra CTACAATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
PhyllanthusFlexu CTACGATTATGCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
PimelodendronGri CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAG-----ATAGTAAAATCTCATAATTTACGATCA  
RicinusCommunis CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAG-----ATAGTAAAATCTCATAATTTACGATCA  
SacoglottisSp CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
ScageaOligostemo CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
TrigoniasBolivia CGACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
AlbiziaTomenosaJ CTACAATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
KerriaJaponica GTACGATTAGTATCTG-----CGACAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
DatiscaCannabina CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA  
QuercusRubra CTACGATTAGTATCTTCT-----TTAGAAAGGTCAGAA-----ATAGTAAAATCTCATAATTTACGATCA





[illegible]









L. perenne TTTCTTCGTAACCAAGTTTTTTATTACGATCAACCTTTTCTAGCTTTCTTCTTGAGCGAAATTTTTTCTAT  
 L. pratense TTTCTTCGTAACCAAGTTTTTTATTACGATCAACCTTTTCTAGCTTTCTTCTTGAGCGAAATTTTTTCTAT  
 L. narbonense TTTATTCGTAACCAAGTTTTTTATTACGATCAACCTTTTCTAGCTTTCTTCTTGAGCGAAATTTGTCTAT  
 C. selaginoides TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 L. hirsutum TTTCTTCGTAACCAATCTTTTATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAATTTTTCTAT  
 L. hypericifolium TTTCTTCGTAACCAATCTTTTATTACGATCAACCTTTTCTAG-----  
 L. usitatissimum TTTCTTCGTAACCAATCTTTTATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAATTTTTTCTAT  
 L. nervosum TTTCTTCGTAACCAAGTTTTTTATTACGATCAACCTTTTCTAGCTTCTTCTTGAGCGAAATTTTTTCTAT  
 L. bienne TTTCTTCGTAACCAAGTTTTTTATTACGATCAACCTTTTCTAGCTTCTTCTTGAGCGAAATTTGTCTAT  
 L. monogynum TTTCTTCGTAACCAAGTTTTTTATTACGATCAACCTTTTCTAGCTTCTTCTTGAGCGAAATTTGTCTAT  
 L. suffruticosum TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 H. clevelandii TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 H. disjunctum TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 L. berlandieri TTTCTTCGTAACCAATCTTTTCATTACGATCCGCTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 L. rupestre TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 D. pentagyna TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 R. indica TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 H. drymarioides TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 H. micranthum TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 H. castaneifolia TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 S. digynum TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 L. oligophyllum TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 H. neblinae TTTCTTCGTAACCAATCTTGTCAATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 H. mystax TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 H. busseana TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 H. gabunensis TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 H. obtusifolia TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 H. neocaledonic TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 H. penicillanth TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 H. spicata TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 H. planchoni TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 H. orientalis TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 H. longipes TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 L. volkensii TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 L. flavum TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTATCAGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 L. catharticum TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 L. arboreum TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 R. linoides TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTATCAGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 R. schomburgkii TTTATTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 R. monsalveae TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 A. pubescens TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 A. khasayama TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 A. saxatilis TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 L. comptonii TTTCTTCGTAACCAATCTTTTCATTACGATCCACTTTTTATCAGTTTCTTCTTGAGCGAAAGTATTTCTAT  
 H. humiriifolia TTTGTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTAGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 I. griffithiana TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTTAT  
 P. magnifolia TTTATTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGGTTCCCTTCTTGAGCGAAAGTATTTCTAT  
 T. sinensis TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGATTCTTCTTGAGCGAAAGTATTTCTAT  
 T. ovoidea TTTCTTCGTAACCAATCTTTTCATTACGATCAACCTTTTCTCGATTCTTCTTGAGCGAAAGTATTTCTAT  
 R. calophylla -----  
 AcalyphaInsulana TTTCTCCGTACCAATCCTTTTCATTACGATCAACATTTTCTCGAGTCTTCTTGAGCGAAATTTATTTCTAT  
 BalanopsPancheri TTTCTCCGTAAACCAATCCTTTTCATTACGATCAACATTTTCTCGGTTCTTCTTGAGCGAAATCTATTTCTAT  
 BischofiaJavanic TTTCTTCGTAAGCAATCTTTTCATTACGATCAACATTTTCTCGGATCTTCTTGAGCGAAATATATTTCTAT  
 BruguieraGymnorh TTTCTCCGTAAACCAATCCTCTTATTTTACGATCAACATTTTCTTGGGTACTTCTTGAGCGAAATATTTCTAT  
 ByrsonimaCrassif TTTCTCCGTAAACCAATCCTTTTATTTCGATCAACATTTTCTCGGAGTCTTCTTGAGCGAAATATATTTCTAT  
 MalpighiaGlabra TTTCTCCGTAAACCAATCCTTTTATTTCGATCAACATTTTCTCGGGTGCTTCTTGAGCGAAATATATTTCTAT  
 CampostylusMann TTTCTTCGTAACCAATCCTTTTCATTACGATCGACATTTTCTCGGTTCTTCTTGAGCGAAATATATTTCTAT  
 ClusiaRosea TTTTTCGTAATCCAATCCTTTTCATTACGATCAACATTTTCTCAGGTTCTTATTGAGCGAAATATATTTTAT  
 CratoxylumCochin TTTCTTCGTAATCCAATCCTTTTCATTACGATCCAACTTTTCTCAGGTTCTTTTCTTGAGCGAAATCTATTTCTAT  
 CtenolophonEngle TTTCTCCGTAAACCAATCCTTTTATTACGATCAACATTTTCTCGAGTTCTTCTTGAGCGAAATATATTTCTAT  
 DrypetesLittoral TTTCTCCGTAAACCAATCCTTTTCATTACGATCAACATTTTCTCGGTTCTTCTTGAGCGAAATATCTTTCTAT  
 ElatineTriandra TTTCTCCGTAAACCAATCCTTTTCATTACGATCAACATTTTCTCGGTTCTTCTTGAGCGAAATATATTTCTAT  
 ErythroxylumNovo TTTCTCCGTAAACCAAGCCTTTTATTACGATCAATATTTTCTCGGTTCTTCTTGAGCGGATAGATTTCTAT  
 EuonymusHamilton TTTCTACGTAAACGAAGAATTTTCATTATGGTCAACATCTCTCGGTTCTTCTTGAGCGAAATCCATTTCTAT  
 EuphorbiaHumifus TTTCTCCGTAAATCAGTGCTTTTCATTACGATCAACATTTTCTCGGTTCTTCTTGAGCGAAATTTTTTCTAT  
 EuphorbiaGuianen TTTCTCCGTAAAGCAATCTTTTCATTATTAATCAACAGTTTCTCGGTTCTTCTTGAGCGAAATTTATTTCTAT  
 FlacourtiaIndica TTTCTTCGTAATCAATCCTTTTCATTTCGATTAATATTTTCTCAGGTTCTTCTTGAGCGAAATATATTTCTAT  
 FlueggeaVirosa TTTCTCCGTAAACCAATCCTTTTCATTACGATCAACATTTTCTCGGTTCTTCTTGAGCGAAATATATTTTAT





L. usitatissimum GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATATATTTTAGCT-----TCAAAGGGT  
 L. nervosum GATACT-----TTCATGCATTATGCTAGATATCAAGGAAAATATATTTTAGCT-----TCAAAGGGT  
 L. bienne GATACT-----TTCATGCATTATGCTAGATATCAAGGAAAATTTTTTTTAGCT-----TCAAAGGGT  
 L. monogynum GATACT-----TTCATGCATTATGCTAGATATCAAGGAAAATTTATTTTAGCT-----TCAAAGGGT  
 L. suffruticosum GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTTAGCT-----TCAAAGGGT  
 H. clevelandii GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTTAGCT-----TCAAAGGGT  
 H. disjunctum GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTTAGCT-----TCAAAGGGT  
 L. berlandieri GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATATATTCTAGCT-----TCAAAGGGT  
 L. rupestre GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTTAGCT-----TCAAAGGGT  
 D. pentagyna GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 R. indica GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATATATTCTAGCT-----TCAAAGGGT  
 H. drymarioides GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 H. micranthum GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTTAGCT-----TCAAAGGGT  
 H. castaneifolia GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 S. digynum GAGCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 L. oligophyllum GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 H. neblinae GATCCT-----TTCGTGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 H. mystax GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 H. busseana GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 H. gabunensis GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 H. obtusifolia GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 H. neocaledonic GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 H. penicillanth GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 H. spicata GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 H. planchoni GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 H. orientalis GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 H. longipes GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 L. volkensii GATCCG-----TTTATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 L. flavum GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 L. catharticum GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 L. arboreum GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 R. linoides GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTTAGCT-----GGAAGGGT  
 R. schomburgkii GATCCT-----TTCGTGCATTATGCTAGATATCAAGGAAAATACATTTCTGGCT-----TCAAAGGGT  
 R. monsalveae GATCCT-----TTCGTGCATTATGCTAGATATCAAGGAAAATACATTTCTGGCT-----TCAAAGGGT  
 A. pubescens GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 A. khasayama GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 A. saxatilis GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 L. comptonii GATCCG-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 H. humiriifolia GATCCT-----TTCGTGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 I. griffithiana GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 P. magnifolia GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACGTTCTGGCT-----TCAAAGGGT  
 T. sinensis GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 T. ovoidea GATCCT-----TTCATGCATTATGCTAGATATCAAGGAAAATACATTTCTAGCT-----TCAAAGGGT  
 R. calophylla -----  
 AcalyphaInsulana GATCCT-----AACATGCATTCTGTTAAATATCAAGGAAAATCCATTTTGGCT-----TCAAAGGAT  
 BalanopsPancheri GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAAGCCATTTCTGGCT-----TCAAAGGAT  
 BischofiaJavanic GATCCT-----TTTATGCATTATGTTAGATATCAAGGAAAAGTTCAATTTGGCT-----TCAAAGGAT  
 BruguieraGymnorh GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAATTAATTTTGGTT-----TCAAAGTAT  
 ByrsonimaCrassif GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAACACATTTAGCT-----TCAAAGGAT  
 MalpighiaGlabra GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAAGGCATTTCTGGCT-----TCAAAGGAT  
 CampostylusMann GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAAGTCAATCTGGCT-----TCAAAGGAT  
 ClusiaRosea GATCCT-----AACATGCATTATGTTAGATATCAAGGAAAATCCATTTTGGCT-----TCAAAGGGT  
 CratogeomysCochin GATCCT-----CGCATGCATTATGTTAGATATCAAGGAAAATTTCTTTTGGCT-----TCAAAGGGT  
 CtenolophonEngle GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAATGCATTTTGGCT-----TCAAAGGAT  
 DrypetesLittoral GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAATCAATTTCTGTCTTCAAATTTCAAAGGGT  
 ElatineTriandra GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAATCCATTTCTGGCT-----TCAAAGGAT  
 ErythroxylumNovo GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAATCCATTTCTGGCT-----TCAAAGGAT  
 EuonymusHamilton GATCCT-----TGCACACATTATGTTAGATATCAAGGAAAATTAATTTCTGGCT-----TCAAAGGAT  
 EuphorbiaHumifus GATCCT-----TTGATGCATTATGTTAGATATCAAGGAAAATCAATTTCTGGCT-----TCAAAGGAT  
 EuphoniaGuianen GATGCT-----TTCATGCATTATGTTAGATATCAAGGAAAATCAATTTCTGGCT-----TCAAAGGAT  
 FlacourtiaIndica GATCTT-----TTCATACATTATGTTAGATATCAAGGAAAATCCATTTTGGCT-----TCAAAGGAT  
 FlueggeaVirosa GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAATCAATTTCTGGCT-----TCAAAGGAT  
 GarciniaSubellip GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAATCAATTTTGGCT-----TCAAAGGAT  
 GomphiaSerrata GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAATCCATTTCTGGCT-----TCAAAGGAT  
 HumiriaBalsamife GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAATCAATTTCTGGCT-----TCAAAGGAT  
 IonidiumCommune GACTCT-----TTCATGCATTATGTTAGATATCAAGGAAAATCAATTTCTGGCT-----TCAAAGGAT  
 IrvingiaMalayana GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAATCAATTTCTGACT-----TCAAAGGAT  
 IxonanthesReticu GATCCT-----TTCATGCATTATGTTAGATATCAAGGAAAATCAATTTCTGACT-----TCAAAGGAT







H. disjunctum TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCACTTTTTAGGCTAT  
L. berlandieri TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCAACTTTTTAGGCTAT  
L. rupestre TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCACTTTTTAGGCTAT  
D. pentagyna TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGATTTTTGGGCTAT  
R. indica TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCACTTTTTAGGCTAT  
H. drymarioides TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAATATTCTATCCACTTTTTAGGCTAT  
H. micranthum TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCACTTTTTAGGCTAT  
H. castaneifolia TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
S. digynum TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCACTTTTTAGGCTAT  
L. oligophyllum TGGTCTCAACCAAAAAACATCTATTAAAACTCA-----TTATACAAACATTCTATCCACTTTTTAGGCTAT  
H. neblinae TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
H. mystax TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
H. busseana TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
H. gabunensis TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
H. obtusifolia TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
H. neocaledonic TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
H. penicillanth TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
H. spicata TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
H. planchoni TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
H. orientalis TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
H. longipes TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
L. volkensii TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACAGTCTATTCACTTTTAGGCTAT  
L. flavum TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCACTTTTTAGGCTAT  
L. catharticum TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCACTTTTTAGGCTAT  
L. arboreum TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCACTTTTTAGGCTAT  
R. linoides TGGTCGCAACCAGAAAAACATTTATTAAAACTCA-----TTATCCAAACATTCTATCCAGTTTTAGGTTAT  
R. schomburgkii TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
R. monsalveae TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
A. pubescens TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCACTTTTTAGGCTAT  
A. khasayama TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCACTTTTTAGGCTAT  
A. saxatilis TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCACTTTTTAGGCTAT  
L. comptonii TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACATTCTATCCGCTTTTTAGGCTAT  
H. humiriifolia TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
I. griffithiana TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCGACTTTTTGGGCTAT  
P. magnifolia TGGTTTCAGCCAGAAAAACATCTATCTAAACTCA-----TTATCCAAGCATTCTATCAACTTTTTGGGCTAT  
T. sinensis TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACAGGATATCCACTTTTTAGGCTAT  
T. ovoidea TGGTCTCAACCAGAAAAACATCTATTAAAACTCA-----TTATCCAAACAGGATATCCACTTTTTAGGCTAT  
R. calophylla -----  
AcalyphaInsulana TGGTTTCAACCAGAAAAGATCTCTATCAATTCA-----TTATCTAAGCATTCTCTCAACTTTTTGGCTAC  
BalanopsPancheri TGGTTTCAACCAGAAAAGGATCAACATAAGCCCA-----TTATACAAGCGCTCTCTCGACTTTCTGTGCTA  
BischofiaJavanica TGGTTTCAACCAGAAAAGGATCTATATAAACCCA-----TTAGCCAAGCATTCTCTCGACTTTTTGGGTTAT  
BruguieraGymnorhiza TGGTTTCAACCAGAAAAGATCTATATAAACCTC-----TTATCCAAGCATTCTCTAGACTTTTTGGGTTAT  
ByrsonimaCrassifolia TGGTTTCATCCAGAACCAATCTATATAAAGTCA-----TTATCCAAGCATTCTCTGACTTTTTGGGTTAT  
MalpighiaGlabra TGGTTTCATCCAGAAATCGATCTATATGAAGTCA-----TTATCCAACCATCTCTTGACTTTTTGGGTTAT  
CamptostylusMann TGGTTTCAACCAGAAAAGATCTATATAAACCCA-----TTATCTAAGCATTCTCTCGGCTTTTTGGGTTCT  
ClusiaRosea TGGTCTCAACCCGACCAGATCTATATAAACCCA-----TTATCTAAGCATTCTCTTGACTTTGTGGCTTAT  
CratogeomysCochin TGGTTTCAACCAGAAAAGATCTATATAAATATA-----TTAGAGAAGCCATCTATCGACTTTATGGGTTAT  
CtenolophonEngle TGGTTTCAACCAGAAAAGGATCTATATAAATCCA-----TTATCCAAGCATTCTCTACCTTTTTGGGCTAT  
DrypetesLittoral TGGTTCAGCCAGAAAAGATCTATATAAACCCA-----TTATCCAAGCATTCTCTCGGCTTTTTGGGCTAT  
ElatineTriandra TGGTTTCATCCAGAAAAATATATATAAAGTCA-----TTAGCCAAGCATTCTCTCGACTTTTTGGGCTAT  
ErythroxylumNovo TGGTTTCAGCCGAAAAGATCTATATAAATCCA-----TTATCCAACCATCTCTAGACTTTTTGGGCTAT  
EuonymusHamilton TGGTCTCAACCAGAAAAGGATCTATATAAACCAA-----TTATCCAAGCATTCTCTCGACTTTTTAGGCTAT  
EuphorbiaHumifusa TGGTTTCAACCAGAAAAGATCTATATCAATTCA-----TTATCCAAACATTCTCTCCAGTTTTGGGCTAT  
EuphoniaGuianensis TGGTTTCAACCAGAAAAGATCTATATAAATCCA-----TTATACAAGCGTCTCTCAACTTTTTGGGCTAT  
FlacourtiaIndica TGGTTTCAACCAGAAAAGATCTATATAAACCCA-----TTATCCAAGCATTCTCTCGACTTTTTGGGCTAT  
FlueggeaVirosa TGGTTTCAACCAGAAAAGATCTATATAAACCCA-----TTATCCAAGCATTCTCTCGACTTTTTGGGCTAT  
GarciniaSubellip TGGTTTCAACCAGAAAAGATCTATATAAATCCA-----TTATCCAAGCATTCTCTCAACTTTTTGGGCTAT  
GomphiaSerrata TGGTTTCAGCCAGAAAAGATCTATATAAATCCA-----TTATCCAAGCATTCTCTCGACTTTTTGGGTTAT  
HumiriaBalsamifera TGGTTTCAACCAGAAAAGATCTATATAAACCCA-----TTATCCAACCATCTCTCGACTTTTTGGGCTAT  
IonidiumCommune GGGTTTGGCCGAAAAGATCTATATAAATCCA-----TTATCCAAGCATTCTCTTGCTTTTTGGGTTAT  
IrvingiaMalayana TGGTTTCAACCAGAAAAGATCTATATAAATCCA-----TTATCCAAGCATTCTCTCGACTTTTTGGGTTAT  
IxanthusReticulatus TGGTTTCAACCAGAAAAGATCTATATAAATCCA-----TTATCCAACCATCTCTCGACTTTTTAGGTTAT  
JatrophaIntegerrima TGGTTTCAACCAGAAAAGATCTATATAAATCCA-----TTATCCAAGCATTCTCTCAACTTTTTGGGCTAT  
KiggelariaAfrica CGGTTTCAATCAAAAAGATCTATATAAACCCA-----TTATCCAACCATCTCTCGACTTATTAGGCTAT  
LacistemaAggregata TGGTTTCAACCAGAAAAGATCTATATAAATCCA-----TTATCCAAGCATTCTCTCGACTTTTTAGGTTAT  
LunaniaParviflora TGGTTTCAACCAGAAAAGATCTATATAAATCCA-----TTATCCAAGCATTCTCTCGACTTTTTAGGTTAT  
ManihotEsculenta TGGTTTCAACCAGAAAAGATCTATATAAATCCA-----TTATCCAAGCATTCTCTCAACTTTTTGGGCTAT  
OxalisCorniculata TGGTCTCAACCAGAAAAGGATCTATATAAACCAA-----TTATCCAAGCATTCTCTCGACTTTTTGGGATAT





H. drymarioides ATGGAT-----AAGACTACGCAGAAATTTGATACAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
H. micranthum ATGGAT-----AATACTACGCAGAAATTTGATACAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
H. castaneifolia ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
S. digynum ATGGAT-----AATACTACGCAGAAATTTGATACAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
L. oligophyllum ATGGAT-----AATACTGCGCAGAAATTTGATACAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
H. neblinae ATAGAT-----AATACTATGAAGAAATGTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
H. mystax ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
H. busseana ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
H. gabunensis ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
H. obtusifolia ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
H. neocaledonic ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
H. penicillanth ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
H. spicata ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
H. planchoni ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
H. orientalis ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
H. longipes ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
L. volkensii ATGGAG-----AATACTACGCAGAAATTTGATACAATAATTCCAATTTATTCGTTGATTGGAACCTTGGAA  
L. flavum ATGGAT-----AATACTACGCAGAAATTTGATACAATAATTCCATTCTTCGTTGATTGGAGCCTTGGAA  
L. catharticum ATGGAT-----AATACTACGCAGAAATTTGATACAATAATTCCAATTTTTCGTTGATTGGAGCCTTGGAA  
L. arboreum ATGGAT-----AATACTACGCAGAAATTTGATACAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
R. linoides ATAGAT-----AATACTACGCATAAATTTGATCCAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
R. schomburgkii GTAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
R. monsalveae GTAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGATTGGATCCTTGGCA  
A. pubescens ATAGAT-----AATACTACGCAGAAATTTGATACAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
A. khasayama ATAGAT-----AATACTACGCAGAAATTTGATACAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
A. saxatilis ATAGAT-----AATACTACGCAGAAATTTGATACAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
L. comptonii ATGGAT-----AATACTACGCAGAAATTTGATACAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
H. humiriifolia -----  
I. griffithiana ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGGGTGGATCCTTAGCA  
P. magnifolia ATAGAT-----AATACTATGAAGAAATTTGATACAATAGTTCCAATTATTCGTTGGGTGGATCCTTAGCA  
T. sinensis ATAGAC-----AATACTACGCAGAAATTTGATACAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
T. ovoidea ATAGAC-----AATACTACGCAGAAATTTGATACAATAATTCCAATTCCTTCGTTGATTGGAGCCTTGGAA  
R. calophylla -----  
AcalyphaInsulana ATAGATAAAGATAAATACTCTGAAGAACTCGATATAATAGTTCCAATTATTCCTTTAATTGGATCATTGGCA  
BalanopsPancheri ATAGAT-----AATAATATGAAGAACTCGATACAATAGTTCCAATTAGTTCCTGTGATTGAATCATTGGCA  
BischofiaJavanica ATAGAT-----AATACTATGAAGAAAGTCGATACAATAGTTCCAATTATGCCTTTGATTGATCATTGGTA  
BruguieraGymnorh CTAGAT-----AATGCTATAAAAAAATCGACACAATAGTTCCAATTATTCCTTTGATTGGATCATTGGCA  
ByrsonimaCrassif ATAGAT-----AATATTATGAAGAACTCGATTCAATAGTTCCAATTATTCCTTTGATTGGATCGTTGGCA  
MalpighiaGlabra ATAGAT-----AATATTAGAAAGAACTTGATTCAAGAGTTCCAATTATTCCTTTGATTGGATCATTGGCA  
CampostylusMann ATAGAT-----AATACTATGAATAAACTCGATACAATAGTTCCAATTATTCCTTTGATTGGATCATTGGCA  
ClusiaRosea ATAGAT-----AATACTATGAAGAACTCGATACAATAGTTCCAATTATTCCTTTGATTGGATCATTGGCA  
CratogeomysCochin ATAGAG-----AATACTATGAAGAACTGGATACAATAGTTCCAATTATTCCTTTGATTGAATCATTGGCA  
CtenolophonEngle ATAGAT-----AATACTCTGAAGAACTCGATACAATAATTCCAATTCCTTTGATTGGATCATTGGCA  
DrypetesLittoral ATAGAC-----AATACTATGAAGAAATCGATACAACAGTTCCAATTATTCCTTTGATTGGATCATTGGCA  
ElatineTriandra ATAGAT-----AATACTAGCAAGAAATTCGAGCCACAGTTCCTATTATTCCTTTGATTGGATCATTGGCA  
ErythroxylumNovo ATAGAT-----AATACTATGAAGAACTCGATTCAACAGTTCCAATTATTCCTTTGATTGGAACATTGGCA  
EuonymusHamilton ATAGAT-----AACGCTATGAAGAACTCGATACAAGAGTTCCACTTATTCTTTGATTGGATCATTGGCA  
EuphorbiaHumifus ATAGATAAAGATAAATACTATGAAGAACTCGATACAATAGTTCCAATTATTCCTTTAATTAGATTATTGGCA  
EuphorbiaGuianen ATAGAT-----AATAATATGAAGAACTTGATACAATAGTTCCAATAATTCCCTGTGATTGAATCATTGGCA  
FlacourtiaIndica ATAGAG-----AATACTATGAATAAACTCGATACAACAGTTCCAATTATTCCTTTGATTGGATCATTAGCA  
FlueggeaVirosa ATAGAT-----AATGCTATGAAGAAAGTTGATACAAGAAATTCCAATTATTCCTTTGATTGGATCATTGGCA  
GarciniaSubellip ATAGAT-----AATACTATGAAGAACTCGATACAATAGTTCCAATTATTCCTTTGATTGGATCATTGTCA  
GomphiaSerrata ATAGAT-----AATACTATGAAGAACTCGATACAATAGTTCCAATCATTCCTTTGATTGGATCCTTGGCA  
HumiriaBalsamife ATAGAT-----AATACTATGAAGAACTCGATACAATAGTTCCAATTATTCCTTTGATTGGATCATTGGCA  
IonidiumCommune ATTGAT-----AACACCATAAAAAAATCGATACAATCGTTCCAATTCTTCCTTTGATTGAATCATTAGAA  
IrvingiaMalayana ATGGAT-----AATACTATGGAGAACTTGATACAATAGTTCCAATTATTCCTTTGATTGGATCATTAGCA  
IxanthosReticu ATAGAT-----AATACTATGAAGAACTCAATACAATAGTTCCAATTATTCCTTTGATTGGATCATTAGCA  
JatrophaIntegerr ATAGATAAAGCTAATACTATGAAGAACTCGATACAATAGTTCCAATTATTCCTTTGATTGGATCATTGTCA  
KiggelariaAfrica ATAGAT-----AATACTATGAATAAACTCGATACAATAGTTCCAATTATTCCTTTGATTGGGTATTGGCA  
LacistemaAggrega ATAGAT-----AATACTATGAATAAACTCGATACAATAGTTCGATATTTCCTTTGATTGGATCATTGGCA  
LunaniaParviflor ATAGAT-----AATACTATGAATAAACTCGATACAATAGTTCCAATTATTCCTTTGATTGGATCATTAGCA  
ManihotEsculenta ATAGATAAAGATAGTACTATGAAGAACTCGATACAATAGTTCCAATTATTCCTTTGATTGGATCATTGGCA  
OxalisCorniculat ATAGAT-----AATGTTATGAAGAACTCGATACAATATTTCCAATTATTCCTCTGTTGGATCATTGTCA  
PassifloraQuadra ATAGAT-----AATATTAGAAAGAACTTGATTCAAGAGTTCCAATTATTCCTTTGATTGGATCATTGTCA  
PhyllanthusFlexu ATAGAT-----AATGCTATGAAGAAAGTTGATACAAGAAATTCCAATTTTTCCTTTGATTGGATCATTGGCA  
PimelodendronGri ATAGATAAAGATAAATACTATGAAGAACTCGATACAATAGTTCCAATTATTCCTTTGATTGGATCATTGGCA  
RicinusCommunis ATAGATAAAGATAAATACTATGAAGAACTCGATACAATAGTTCCAATTATTCCTTTAATTAGATCATTGGCA  
SacoglottisSp ATAGAT-----AATACTATGAAGAACTCGATACAATAGTTCCAATTATTCCTTTGATTGGATCATTGGCA



ByrsonimaCrassif	AAAACAAAATTTTGTAAACGCAGTAGGACATCCCATTAGTAAACCGAGCTGGGCGGAT
MalpighiaGlabra	AAAACAAAATTTTGTAAACGCAGTAGGGCATCCCATTAGTAAACCGAGCTGGGCGGAT
CamptostylusMann	AAAACGAAATTTTGTAAACGAGCGGGACATCCCATTAGCAAACCGGCTCGGGCGGAT
ClusiaRosea	AAAATGAAATTTTGTAAATGCAGTAGGACATTCTGTAGTAAATCGACCTGGGCTGAT
CratoxylumCochin	AAAATGAAATTTTGTAAACGAAGTAGGACATCCCATTAGTAAACCGACCTGGACCGAT
CtenolophonEngle	AAAAGAAAATTTTGTAAATGCAGTAGGGCATCCCATTAGTAAACTGACCCGGGCTGAT
DrypetesLittoral	AAAACGAAATTTTGTAAACGCAGTGGGGCATCCCATTAGTAAACCGACCCGGGCCGAT
ElatineTriandra	AAAACGAAATTTTGTAAACACAACAGGACAGCCCATTAGTAAAGCGAGCTGGGCTGAT
ErythroxylumNovo	AAAACGAGATTTTGTAAACGAAAAGGACATCCCATTAGTAACTCGGCCCGGGCTGAT
EuonymusHamilton	AAAGCGAGATTTTGTAAACGTATTAGGACATCCCATTAGTAAACGACCCGGGCTGAT
EuphorbiaHumifus	AAAACGAAATTTTGTAAACGCAGTAGGACATCCTATTAGTAAACCGATCCGGACTCAT
EuphroniaGuianen	AAAACAAAATTTTGTAAATGTGGTAGGACATCCCCTTAGTAAATCGGACCCGGACTGAT
FlacourtiaIndica	AAAATGAAATTTTGTAAATGCAGTAGGACATCCCATTAGTAAACCGACCTGGGGCGAT
FlueggeaVirosa	AAAACGAAATTTTGTAAATGCAGTAGGACATCCTATTAGTAAACCTATTTGGGCTGGT
GarciniaSubellip	AAAATGACATTTTGTAAATGCAGTAGGATATCCCATTAGTAAATCGACCTGGGCTGAT
GomphiaSerrata	AAAATGAAATTTTGTAAACAGTAGGACATCCCATTAGTAAATCCGACCCGGGCTGGT
HumiriaBalsamife	AAAATGAAATTTTGTAAACGCAGTAGGGCATCCCATTAGTAAACCGATCCGGGCTGAT
IonidiumCommune	AAAATGAAATTTTGTAAATGGAATAGGACAGCCCATTGGTAAACCACTTGGGCTGAT
IrvingiaMalayana	AAAACGAAATTTTGTAAATGCAGTAGGACATCCTATTAGTAAACCGATCCGGGCTGAT
IxonanthesReticu	AAAACGAAATTTTGTAAACGAATAGGCCATCCCATTAGTAAACCGCCTTGGGCTGAT
JatrophaIntegerr	AAAACGAAATTTTGTAAAGGCAGTAGGACATCCCATTAGTAAACCGGTCCGGACTGAT
KiggelariaAfrica	AAAGCAAAATTTTGTAAACGAACAGGACATCCCTTTAGTAAACCGACTTGGGCGGAT
LacistemaAggrega	AAAACGAAATTTTGTAAACGCAGTAGGACATCCCATTAGTAAACCGGCTTGGGCAGAT
LunaniaParviflor	AAAATGAAATTTTGTAAACGTAGTAGGACATCCCATTAGTAAACCGACCTGGGCCGAT
ManihotEsculenta	AAAACGAAATTTTGTAAACGCAGTAGGACATCCCATTAGTAAACCGATTCGGGCGGAT
OxalisCorniculat	AAATCAAAATTTTGTAAACACATTAGGGTATCCCATTAGTAAAGCCTACTTGGGCGGAT
PassifloraQuadra	AAAACAAAATTTTGTAAACGCAGTAGGGCATCCCATTAGTAAACCGAGCTGGGCGGAT
PhyllanthusFlexu	AAAACGAAATTTTGTAAACGCAGTAGGGTATCCTATTAGTAAACCTATTTGGGCTGGT
PimelodendronGri	AAAACGCAATTTTGTAAATGCAGTAGGACATCCCATTAGTAAACCGACCCGGGCTGAT
RicinusCommunis	AAAATGAAATTTTGTAAACGCAGCAGGACATCCCATTAGTAAACCGACCTGGGCGGAT
SacoglottisSp	AAAACGAAATTTTGTAAACGCAGTAGGGCATCCCATTAGTAAACCGACCCGGGCTGAT
ScageaOligostemo	AAAACACAATTTTGTAAACGTAGTAGGACATCCCATTAGTAAACCGATTTGGGCTGAT
TrigoniasBolivian	AAAGCAAAATTTTGTAAACGAACAGGACATCCCTTTAGTAAACCGACTTGGGCGGAT
AlbiziaTomenosaJ	AAAGCAAAATTTTGTAAATGTATTAGGACATCCCATTAGTAAAGCCGGTCTGGGCCGAT
KerriaJaponica	AAAGTTAAATTTTGTAAACGATTAGGACATCCTATTAGTAAAGTCCACCTGGTCCGAT
DatiscaCannabina	AAAGTGCAATTTTGTAAACGTATGGGCGCATCCTATTAGTAAAGTCCACCTGGGCGGAT
QuercusRubra	AAAGCGAAATTTTGTAAACGCAGTAGGGCATCCAATTAGTAAAGCTGACTCGGGCCGAT

## Appendix B: Voucher specimen information for Linaceae accessions utilized in Chapters 3 and 4.

Table A1: Species sampled for phylogenetic analyses, with biogeographic distributions, morphological character states, DNA accession numbers, voucher specimen information, and GenBank accession numbers for previously-published sequences used in these analyses. Distribution categories: eSAM = eastern South America, wSAM = western South America, nNAM = North America North of Mexico, MEX = Mexico, SAF = South Africa, Eur = Eurasia, TAF = tropical Africa. Morphology: fruit segments (0 = ten, 1 = six, 2 = five, 3 = four), pollen type [0 = tricolpate, 1 = few-porate (<12), 2 = multiporate (12-20)], false septa (0 = incomplete, 1 = intermediate, 2 = complete), styler fusion (0 = free, 1 = fused). Accessions from which ITS was cloned are indicated by ‡. Accessions whose sequences were included in the reduced data matrix in Chapter 5 are indicated by \*. Sequences newly obtained for this study are indicated by “-” rather than GenBank accession numbers, which will be obtained at the time of journal publication.

Taxon	Geographic Distribution Category	Morph.	Acc. #	Voucher	Locality	Herbarium	matK, trnK, ndhF, trnL-F, ITS
<i>Cliococca selaginoides</i> (Lam.) Rogers & Mildner	eSAM+wSAM	0,0,2,0	049	Krapovickas & Cristóbal 34169	Brazil, Rio Grande do Sul	MO 2777516	-, -, FJ160774, FJ160858, FJ169540
			206‡ *	Baeza & Rodriguez 1900	Chile, Region VIII, Prov. Arauco	CONC	-, -, -, -
			220‡	JRM 2004-XII-11.1	Uruguay, Cerro Largo	TEX	-, -, -, -
<i>Hesperolinon clevelandii</i> Small	nNAM	1,0,0,0	022*	Ertter 8537	U.S.A., CA, Solano Co.	TEX	-, -, -, -
<i>Hesperolinon disjunctum</i> Sharsmith	nNAM	1,0,0,0	025*	J. McDill 087	U.S.A., CA, San Benito Co.	TEX	-, -, -, -
<i>Hesperolinon drymarioides</i> Small	nNAM	1,0,0,0	047‡ *	Dibble and Griggs 14	U.S.A., CA, Lake Co.	MO 3204309	-, -, -, -

<i>Hesperolinon micranthum</i> Small	nNAM	1,0,0,0	048*	van der Werff & Clark 4574	U.S.A., CA, San Diego	MO 3099280	- ,FJ160818,FJ160775, FJ160859,FJ169542
<i>Linum acuticarpum</i> C.M.Rogers	SAF	0,0,0,1	SA3 †*	McDill 2005-XII-23.1	South Africa, Western Cape	TEX	-,-,-,-,-
<i>Linum adustum</i> E.Mey. ex Planch.	SAF	0,0,0,1	SA1 3	Pillans 7354	South Africa, Western Cape	UCT	-,-,-,-,-
			SA8 †	J. McDill 2006-I-12.1	South Africa, Western Cape, Stilbaai	TEX	-,-,-,-,-
<i>Linum aethiopicum</i> Thunb.	SAF	0,0,0,1	055†	Bremer 577	South Africa, Eastern Cape	MO 4044959	-,-,-,-,-
			SA1 1†	J. McDill 2006-I-13.1	South Africa, Western Cape, Cape Agulhas	TEX	-,-,-,-,-
			SA1 2†*	J. McDill 2006-I-9.1	South Africa, Eastern Cape, Cape St. Francis	TEX	-,-,-,-,-
			SA7 †	J. McDill 2006-I-12.2	South Africa, Western Cape, DeHoop N.P.	TEX	-,-,-,-,-
<i>Linum africanum</i> L.	SAF	0,0,0,1	SA1 *	J. McDill 2005-XII-16.1	South Africa, Western Cape	TEX	-,-,-,-,-
<i>Linum alatum</i> (Small) Winkler	SAF	2,2,2,1	307*	Rogers 13131	U.S.A., TX, Refugio Co.	WUD	-,-,-,-,-
<i>Linum arboreum</i> L.	EUR	0,0,0,0	164*	R. Dechamps 9103	Greece, Crete	MO 3919917	-,-,-,-,-
<i>Linum arenicola</i> (Small) Winkler	nNAM	0,0,1,0	268*	Rogers 12148	U.S.A., FL, Monroe Co.	WUD	-,-,-,-,-
<i>Linum aristatum</i> Engelm.	nNAM	2,2,2,1	041*	J. Spencer & D.	Mexico, Chihuahua	TEX	-,-,-,-,-



				Atwood 536			
			233	J. McDill 2004-VI- 12.1	U.S.A., TX, Loving Co.	TEX	-, -, -, -
<i>Linum australe</i> Heller	nNAM +MEX	2,2,2,1	232	J. McDill 23	U.S.A., AZ, Coconino	TEX	-, -, -, -
<i>Linum bahamense</i> North. ssp. <i>bahamense</i>	nNAM	0,0,0,0	006	D.S. Correll 42766	Bahamas, Little Abaco Isl.	TEX	-, -, -, -
			007*	D.J. Harvey 7746	Bahamas, North Andros Isl.	TEX	-, -, -, -
<i>Linum bahamense</i> North. ssp. <i>corallicola</i> (Small) Rogers	nNAM	0,0,0,0	F1	Correll 44802	Bahamas, Grand Bahama Isl.	F	-, -, -, -
<i>Linum berlandieri</i> ssp. <i>berlandieri</i> Hooker	nNAM	2,2,2,1	027	B.L. Turner 21-427	U.S.A., TX, Kerr Co.	TEX	-, -, -, -
			234	J. McDill 2005-V- 14.1	U.S.A., TX, Randall Co.	TEX	-, -, -, -
			236*	J. McDill 2005-V- 14.4	U.S.A., TX, Potter Co.	TEX	-, -, -, -
			241	JRM 2005-V- 14.5	U.S.A., TX, Hutchins on Co.	TEX	-, -, -, -
			246	J. McDill ??	U.S.A., TX, Kerr Co.	TEX	-, -, -, -
<i>Linum berlandieri</i> ssp. <i>filifolium</i> (Shinners) Rogers	nNAM	2,2,2,1	028*	B.L. Turner 99-127	U.S.A., TX, Val Verde Co.	TEX	-, -, -, -
			291	Sandra Aguilar Ruiz 19	Mexico, Coahuila	TEX	-, -, -, -
<i>Linum bienne</i> Mill.	EUR	n.a.	214	J. McDill 2004- XII-05-1	Chile, Reg. VIII Biobío	TEX	-, -, -, -
<i>Linum brevifolium</i> A.St.-Hil. & Naudin	eSAM	0,0,1,0	193*	R.Mildne r 3	Brazil, Santa Catarina	WUD	-, -, -, -

<i>Linum burkartii</i> Mildner	eSAM	0,0,1,0	221*	JRM 2004-XII-13.1	Uruguay, Paysandú	TEX	-,-,-,-,-
<i>Linum carneum</i> A.St.-Hil.	eSAM	0,0,1,0	200*	Marchesi & Vignale 24-II-1996	Uruguay, Rio Negro	MVFA 25419	-,-,-,-,-
<i>Linum carteri</i> Small	nNAM	2,2,2,1	266*	Rogers 12153	U.S.A., FL, Dade Co.	WUD	-,-,-,-,-
<i>Linum catharticum</i> L.	EUR	0,0,0,0	151	Skvortsov et al. 4263078	Estonia	MO 4263078	-,-,-,-,-
			294*	O. Dollenz 682	Chile	GH	-,-,-,-,-
			L43	n.a.	Germany	MJG 040944	- ,FJ160835,FJ160796, FJ160880,FJ169533
<i>Linum chamissonis</i> Schiede	wSAM	0,0,1,0	195*	R. Mildner 15	Chile, Reg. VIII Biobío	WUD	-,-,-,-,-
			215*	JRM 2004-XII-05.2	Chile, Reg. VIII Biobío	TEX	-,-,-,-,-
<i>Linum compactum</i> A. Nelson	nNAM	2,2,2,1	231	B.B. Simpson 1-V-05-1	U.S.A., TX, Colorado Co.	TEX	-,-,-,-,-
			SD1 *	J. McDill 2008-VII-20.1	U.S.A., SD, Jackson Co.	TEX	-,-,-,-,-
<i>Linum comptonii</i> C.M.Rogers	SAF	0,0,0,1	059*	Esterhuysen 35882	South Africa, Western Cape, Stellenbosch	MO	- ,FJ160821,FJ160778, FJ160862,FJ169550
			SA1 9	Esterhuysen 35882	South Africa, Western Cape, Stellenbosch	UCT	-,-,-,-,-
<i>Linum cremnophilum</i> I.M.Johnst.	wSAM	0,0,0,1	187*	J.M. Johnston 5707	Chile, Reg. II Antofagasta	WUD	-,-,-,-,-
<i>Linum cruciata</i> Planch.	MEX	0,0,0,0	F2*	H.S. Gentry 7194	Mexico, Sinaloa	F	-,-,-,-,-
<i>Linum decumbens</i> Desf.	EUR	n.a.	162	A. Dubuis s.n.	Algeria	MO 4359542	-,-,-,-,-

<i>Linum elegans</i> Spruner ex Boiss.	EUR	0,0,0,0	163	C. Evrard 10.826	Greece	MO 3919889	-,-,-,-,-
<i>Linum elongatum</i> (Small) Winkler	nNAM	2,2,2,1	253*	J. McDill 30-III-03- 3	U.S.A., TX, La Salle Co.	TEX	-,-,-,-,-
<i>Linum erigeroides</i> A.St.-Hil.	eSAM	0,0,1,0	196	Marchesi & Vignale 24-X- 1997	Uruguay, Rio Negro	MVFA	-,-,-,-,-
<i>Linum esterhuysenae</i> C.M.Rogers	SAF	0,0,0,1	061	Rogers 13695	South Africa, Western Cape, Swartber g Pass	MO 3671760	-, FJ160822, FJ160779, FJ160863, FJ169549
			SA9 *	J. McDill SA9	South Africa, Western Cape, Swartber g Pass	TEX	-,-,-,-,-
			SA1 0	J. McDill SA10	South Africa, Western Cape, Swartber g Pass	TEX	-,-,-,-,-
<i>Linum flagellare</i> (Small) Winkler	MEX	0,0,0,0/ 1	042*	Guy Nesom 5359	Mexico, Nuevo Leon	TEX	-,-,-,-,-
<i>Linum flavum</i> L.	EUR	0,0,0,0	149*	C.M. Rogers 13374	Turkey	MO 2372238	-,-,-,-,-
			L57	H. Weber	Romania	MJG R00024	-, FJ160833, FJ160794, FJ160878, FJ169538
<i>Linum floridanum</i> (Planch) Trel. var. <i>chrysocarpum</i> Rogers	nNAM	0,1,1,0	124*	Loran C. Anderson 10897	U.S.A., FL, Franklin Co.	MO 3666555	-,-,-,-,-
<i>Linum gracile</i> Planch.	SAF	0,0,0,1	SA4	J. McDill 2005- XII-22.1	South Africa, Western Cape, Montagu Pass	TEX	-,-,-,-,-
			SA5	J. McDill 2005- XII-31.1	South Africa, Western Cape, Tradouw	TEX	-,-,-,-,-

					Pass		
			SA6*	J. McDill 2006-I- 10.1	South Africa, Western Cape, Grootrivi er Pass	TEX	-,-,-,-,-
<i>Linum guatemalense</i> Benth.	MEX	0,0,1,0	293*	Proctor 25354		TEX	-,-,-,-,-
<i>Linum heterostylum</i> C.M.Rogers	SAF	0,0,0,1	062	C.M. Rogers 13694	South Africa, Western Cape	MO	-,-,-,-,-
<i>Linum holstii</i> Engl. ex R.Wilczek	TAF	0,0,0,0	139*	LaCroix 3897	Malawi	MO	-,-,-,-,-
<i>Linum hudsonioides</i> Planch.	nNAM	2,2,2,1	032	W.R. Carr 14755	U.S.A., TX, Brown Co.	TEX	-,-,-,-,-
			244	J. McDill 2004-V- 7.3	U.S.A., TX, Blanco Co.	TEX	-,-,-,-,-
			250*	Hillis 141	U.S.A., TX, Mason Co.	TEX	-,-,-,-,-
<i>Linum hypericifolium</i> Salisb.	EUR	n.a.	148	M. Merello et al. 2116	Republic of Georgia	MO 04973936	-, FJ160828, FJ160789, FJ160873, -
<i>Linum imbricatum</i> (Raf.) Shinnery	nNAM	2,2,2,1	005	W.R. Carr 18038	U.S.A., TX, Brazoria Co.	TEX	-,-,-,-,-
			249*	J. McDill 2002-V- 22.1	U.S.A., TX, Travis Co., Brackenri dge Field Lab	TEX	-,-,-,-,-
<i>Linum intercursum</i> Bicknell	nNAM	0,1,0,0	401	Rogers 13194	U.S.A., NC, Davidson Co.	WUD	-,-,-,-,-
<i>Linum kingii</i> S. Watson	nNAM	0,0,0,0	123*	M.D. Windham	U.S.A., UT,	MO	-,-,-,-,-

				92-100	Garfield Co.		
			223	J. McDill 021	U.S.A., UT, Salt Lake Co.	TEX	-,-,-,-,-
<i>Linum lasiocarpum</i> Rose	MEX	0,0,0,0	310	Rogers 13455	Mexico, Nuevo Leon	WUD	-,-,-,-,-
<i>Linum lewisii</i> Pursh	nNAM	n.a.	010	B.L. Turner 21-263	U.S.A., TX, Reeves Co.	TEX	-, FJ160839, FJ160800, FJ160884, -
<i>Linum littorale</i> A.St.-Hil.	eSAM	0,0,1,0	089*	R.M. Harley 26923	Brasil, Bahia	MO 04893852	-,-,-,-,-
<i>Linum longipes</i> Rose	MEX	0,0,0,1	038*	Panero 5302	Mexico, Puebla	TEX	-,-,-,-,-
			040	Breedlove 65221	Mexico, Puebla	TEX	-,-,-,-,-
<i>Linum lundellii</i> Rogers	nNAM	2,2,2,1	186*	D.S. Correll 32270	U.S.A., TX, Starr Co.	TEX	-,-,-,-,-
<i>Linum macraei</i> Benth.	wSAM	0,0,0,1	210*	J. McDill 2004-XII-02.1	Chile, Reg. IV Coquimb o	TEX	-,-,-,-,-
<i>Linum marginale</i> A. Cunn.	Australia	n.a.	076	R. Taylor 127	Australia, South Australia	MO 05023280	-, FJ160843, FJ160804, FJ160888, -
<i>Linum maritimum</i> L.	EUR	0,0,0,0	L29 *	n.a.	Italy	MJG 040937	-, FJ160850, FJ160811, FJ160895, FJ169535
<i>Linum medium</i> (Planch.) Britton	nNAM	0,0,1,0	001	V.E. McNeilus 98-404	U.S.A., TN, Cumberland Co.	TEX	-,-,-,-,-
			256*	J. McDill 2005-V-11.2	U.S.A., TX, Lee Co., Lake Somerv ille	TEX	-,-,-,-,-
			257	JRM 2005-V-11.1	U.S.A., TX, Lee Co., Co. Rd. 309	TEX	-,-,-,-,-
<i>Linum mexicanum</i> HBK	MEX	0,0,0,1	037*	Carmelino Santiz Ruiz 209	Mexico, Chiapas	TEX	-,-,-,-,-
<i>Linum monogynum</i> G. Forst.	Australia	n.a.	224	H. Meudt 245	New Zealand	TEX	-,-,-,-,-

<i>Linum narbonense</i> Desf.	EUR	n.a.	013	C. Ardo et al. 2303	Spain	TEX	-,-,-,-,-
<i>Linum nelsonii</i> Rose	MEX	0,0,0,0	F5	Leavenworth 799	Mexico, Nuevo Leon	F	-,-,-,-,-
<i>Linum neomexicanum</i> Greene	nNAM	0,0,0,0	228*	Springer 3A.6	U.S.A., NM	TEX	-,-,-,-,-
<i>Linum nervosum</i> Waldst. & Kit.	EUR	n.a.	146	n.a.	Russia	MO 4250021	-,-,-,-,-
<i>Linum oligophyllum</i> Willd. ex Schult.	wSAM	0,0,1,0	091*	D.N. Smith et al. 9426	Peru, Ancash	MO 3277219	-, -, FJ160783, FJ160867, FJ169546
<i>Linum orientale</i> Boiss.	EUR	0,0,0,0	143*	n.a.	Russia	MO 3144852	-,-,-,-,-
<i>Linum orizabae</i> Planch.	MEX	0,0,1,1	270*	Itlis et al. 261	Mexico, Morelos	WUD	-,-,-,-,-
<i>Linum pratense</i> Small	nNAM	n.a.	238	McDill 2005-V-13.1	U.S.A., TX, Montague Co.	TEX	-,-,-,-,-
<i>Linum pringlei</i> S. Watson	MEX	0,0,0,0	201*	Henricks on 24148	Mexico	TEX	-,-,-,-,-
<i>Linum prostratum</i> Domb. ex Lam.	wSAM	0,0,0,0	194*	R. Mildner 52	Peru, Lima	WUD	-, -, FJ160784, FJ160868, FJ169545
<i>Linum puberulum</i> (Engelmann) Heller	nNAM +MEX	2,2,2,1	183	A.M. Powell 1881	U.S.A., TX, Ward Co.	TEX	-,-,-,-,-
<i>Linum quadrifolium</i> Thunb.	SAF	0,0,0,0	258*	Rogers 13703	South Africa, Western Cape	WUD	-,-,-,-,-
<i>Linum rigidum</i> Pursh.	nNAM	2,2,2,1	035*	W.R. Carr 18225	U.S.A., TX, Randall Co.	TEX	-,-,-,-,-
			400	McDill 2005-V-15.3	U.S.A., TX, Roberts Co.	TEX	-,-,-,-,-
<i>Linum rupestre</i> (A. Gray) Engelmann	nNAM +MEX	0,0,0,0	034	J. Saunders 1343	U.S.A., TX, Travis Co.	TEX	-,-,-,-,-
			036*	B.L. Turner 21-589	U.S.A., TX, Sutton Co.	TEX	-, FJ160824, FJ160785, FJ160869, FJ169553
			242	J. McDill 2004-8	U.S.A., TX, Blanco	TEX	-,-,-,-,-

					Co.		
<i>Linum rzedowskii</i> Arreguin	MEX	n.a.	165	E. Ventura V. 1051	Mexico	MO 3909940	-, -, -, -
<i>Linum scabrellum</i> Planch.	MEX	0,0,1,0	F7*	Purpus 4923	Mexico, San Luis Potosi	F	-, -, -, -
<i>Linum schiedeanum</i> Schlecht. & Cham.	MEX	0,0,0,0	009*	Alonso Mendez Ton 9708	Mexico, Chiapas	TEX	-, -, -, -
<i>Linum scoparium</i> Griseb.	eSAM	0,0,0,0	219*	JRM 2004-XII-12.1	Uruguay, Rivera	TEX	-, -, -, -
<i>Linum striatum</i> Walter	nNAM	0,0,1,0	075*	A.W. Cusick 34532	U.S.A., OH, Portage Co.	MO 04881564	-, -, -, -
<i>Linum strictum</i> L.	EUR	0,0,0,0	012*	J.L. Panero 6966	Spain, Canary Islands, Tenerife	TEX	-, -, -, -
<i>Linum subteres</i> (Trel.) Winkler	nNAM	2,1,1,1	302	Rogers 13244	U.S.A., UT, San Juan Co.	WUD	-, -, -, -
<i>Linum suffruticosum</i> L.	EUR	0,0,0,0	011*	C. Navarro et al. CN 2339	Spain	TEX	-, FJ160846, FJ160807, FJ160891, FJ169532
<i>Linum sulcatum</i> Ridell	nNAM	0,1,0,1	132*	Henderson 93-207	U.S.A., MO, Bates Co.	MO 4375070	-, -, -, -
<i>Linum tauricum</i> Willd.	EUR	0,0,0,0	157	P. Frost-Olsen 1070	Bulgaria	MO 04642915	-, -, -, -
<i>Linum tenellum</i> Schlecht. & Cham.	MEX	0,0,1,0	F6*	Guy Nesom 6260	Mexico, Tamaulipas	F	-, -, -, -
<i>Linum tenue</i> Desf.	EUR	0,0,0,0	66*	Maroc 12600	Morocco	MO	-, -, -, -
<i>Linum tenuifolium</i> L.	EUR	0,0,0,0	156*	P. Frost-Olsen 2115	Serbia	MO	-, -, -, -
			L17	M. Kropf	Germany	MJG 040934	-, FJ160848, FJ160809, FJ160893, FJ169529
<i>Linum thesioides</i> Bartl.	SAF	0,0,0,0	SA17*	Esterhuysen 4456	South Africa, Western Cape	UCT	-, -, -, -
<i>Linum thunbergii</i> Eckl. & Zeyh.	SAF	0,0,0,0	067‡	Bourell 2728	South Africa, Transvaal	MO 4389048	-, -, -, -

<i>Linum trigynum</i> L.	EUR	0,0,0,0	72*	A.E. Wright 2600	New Zealand	MO 2695749	-, -, -, -
<i>Linum usitatissimum</i> L.	EUR	n.a.	L4	n.a.	cultivated	MJG 040924	-, FJ160842, FJ160803, FJ160887, -
<i>Linum vernale</i> Wooton	nNAM	2,1,1,1	184*	Correll & Correll 38582	U.S.A., TX, El Paso Co.	TEX	-, FJ160851, FJ160812, FJ160896, FJ169552
<i>Linum villosum</i> C.M.Rogers	SAF	0,0,0,0	SA14*	Esterhuysen 27360	South Africa, Western Cape	UCT	-, -, -, -
<i>Linum virginianum</i> L.	SAF	0,0,1,0	130*	R. Kral 53267	U.S.A., AL, Chilton Co.		-, -, -, -
<i>Linum viscosum</i> L.	EUR	n.a.	L41	n.a.		MJG 040939	-, FJ160830, FJ160791, FJ160875, -
<i>Linum volkensii</i> Engl.	TAF	0,0,0,0	137*	Gereau & Kayombo 4662	Tanzania	MO	-, FJ160852, FJ160813, FJ160897, FJ169531
<i>Linum westii</i> Rogers	nNAM	0,1,2,0	269*	S. McDaniel 6560	U.S.A., FL, Jackson Co.		-, -, -, -
<i>Linum</i> sp. nov.	nNAM	2,2,2,1	n.a.*	Sivinski 6843	U.S.A., NM, Eddy Co.	TEX	-, -, -, -
<i>Radiola linoides</i> Roth.	EUR	3,0,0,0	171*	Egeröd 6935	Sweden, Goteborg	MO 3928062	-, -, -, -
<i>Reinwardtia indica</i> Dum.	Southeast Asia	n.a.	044	J. McDill s.n.	Cultivated	TEX	-, FJ160853, FJ160814, FJ160898, -
<i>Sclerolinon digynum</i> (A. Gray) C.M. Rogers	nNAM	3,0,2,0	083	M.S. Taylor 3935	U.S.A., CA, Plumas Co.	MO 3209887	-, FJ160826, FJ160787, FJ160871, FJ169541
			PM1983*	Patterson & Melonghi 1983	U.S.A., CA, Sierra Co.	TEX	-, -, -, -



## Appendix C: Chloroplast Data Alignment, Chapter 3

*matK* (1-1154), *trnK* 3' intron (1155-1560),  
*ndhF* (1561-1994), *trnL-F* (1995-2748)

Taxon/Node	111111111122222222223333333333444444444455555555556666666666777
	123456789012345678901234567890123456789012345678901234567890123456789012
001Lstriatum	AATGAAAAATATAAAAAA?AGAAGTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
005Limbricatum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
006Lbahamense	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
007Lbahamense	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
009Lschiedeanum	AATGAAAAATATAAAAACTATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
010Llewisii	-----
011Lsuffruticosu	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
012Lstrictum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
013Lnarbonense	AAAAGAACAATATAAAAAATATATAG-----GATCCCTTAAAAAGCTTCCTATACCCACTTATCTT
022Hclevelandii	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
025Hdisjunctum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
027Lberlandierib	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
028Lberlandierif	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
032Lhudsonioides	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
034Lrupestre	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
035Lrigidumrigid	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
036Lrupestre	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
037Lmexicanum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
038Llongipes	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
040Lcruciata	---TGAAAAATA-----ATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
041Laristatum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
042Lflagellare	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
044Reinwardtia	AATGGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
047Hdrymarioides	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
048Hmicranthum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
049Cselaginoides	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
055Laethiopicum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
059Lcomptonii	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
061Llesterhuysena	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
066Ltenue	AATGACAAAAATACAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
062Lheterostylum	-----
067Lthunbergii	-----
072Ltrigynum	AATGACAAAAATACAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
075Lstriatum	--TGAAAAATATAAAAAAATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
076Lmarginale	AAAAGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
083Sdigynum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
089Llittorale	AATGCAAAAAATATAAAAAAGATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
091Lloliophyllum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
123Lkingii	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
124Lfloridanum	AATGAAAAATATAAAAAAATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
130Lvirginianum	AATGAAAAATATAAAAAAATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
132Lsulcatum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
137Lvolkensisii	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCGCTTATCTT
139Lholstii	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCGCTTATCTT
143Lorientale	-----AATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
146Lnervosum	AAAGGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
148Lhypericifoli	AATGGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
149Lflavum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
151Lcatharticum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
156Ltenuifolium	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
157Ltauricum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
163Lelegans	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
164Larboreum	AATGAAAAATATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT
165Lrzedowskii	-----

171Rlinoides AATGGAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
183Lpuberulum AATGCAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
184Lvernale AATGCAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
186Llundellii AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
187Lcremophyllu AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
193Lbrevifolium AATGCAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
194Lprostratum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
195Lchamissonis AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
196Lerigeroides -----  
200Lcarneum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
201Lpringlei AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
206Cselaginoides AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
210bLmacraeimacr AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
215Lchamissonis AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
219Lscoparium AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
220Cselaginoides AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
221Lburkartii AATGCAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
223Lkingii A-TGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
224Lmonogynum AAAAGAAAAATAATAAATATTTACAACGGATAGATCCCTTAAAAAGCTTCCTATACCCACTTATCTT  
228Lneomexicanum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
231Lcompactum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
232Laustrale AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
233Laristatum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
234Lrigidumberla AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
236Lrigidumberla AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
238Lpratense -----TATAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
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242Lrueprestre AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
246LrigidumKerr AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
253Lelongatum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
256Lmedium AATGAAAAATAATAAATAATTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
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258Lquadrifolium AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
259Lheterostylum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
266Lcarteri AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
268Larenicola AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
269Lwestii AATGAAAAATAATAAATAATTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
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293Lquatemalense AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
294Lcatharticum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
302Lsubteres AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
307Lalatum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
310Llasiocarpum AATGAAAAATAAAAACTATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
244Lhudsonioides AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
214Lbienne AAAAGAAAAATAATAAATATTTACAACGGATAGATCCCTTAAAAAGCTTCCTATACCCACTTATCTT  
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AlLrigidumpuberu AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
F1Lbahamensecora AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
F2Lcruciata AATGAAAAATAAAAACTATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
F5Lnelsonii AATGAAAAATAAAAACTATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
F6Ltenellum AATGAAAAATAAAAACTATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
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L29Lmaritimum AATGAAAAATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
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L41Lviscosum AATGGAAAAATAAAAAATATTTGCAACTAGATAGATCTCTTCAAAAAAGCTTCCTATACCCACTTATCTT  
L43Lcatharticum AATGAAAAATAAAAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
L57Lflavum -----  
LPB2Loligophyllu AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
PM1983Sdigynum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
SA1Lafricanum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
SA10Lesterhuysen AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
SA11LinumAgulhas AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
SA12Laethiopicum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
SA13LinumspWindb AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT  
SA14Lvillosum AATGAAAAATAATAAATATTTAGAACTAGATGGATCTCTTAAAAAGCTTCCTATACCCACTTATCTT



151Lcatharticum TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGATCCCTTTTTTCGGAAAATCT  
156Ltenuifolium TCGGGAGTATATTTATATAGTTACTTACGTTTCGTGGTTTAAATGGATATGGATCCATTTTTTCAGAAAATGT  
157Ltauricum TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATCGATATGGATCCATTTTTTCGGAACATGT  
163Lelegans TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATCGATATGGATCCATTTTTTCGGAACATGT  
164Larbareum TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATCGATATGGATCCATTTTTTCGGAACATGT  
165Lrzedowskii -----AAGGT  
171Lrinoides TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGATCCATT-----AAGGT  
183Lpuberulum TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGCTCCATTTTTTCAGAAAATGT  
184Lvernale TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGCTCCATTTTTTCAGAAAATGT  
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187Lcremnophyllu TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
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194Lprostratum TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
195Lchamissonis TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
196Lerigeroides -----  
200Lcarneum TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
201Lpringlei TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATAGATATGGCTCCATTTTTTCAGAAAATGT  
206Cselaginoides TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
210bLmacraeimacr TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
215Lchamissonis TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
219Lscoparium TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
220Cselaginoides TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
221Lburkartii TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
223Lkingii TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
224Lmonogynum TCGGGAGTATATTTATATAGTTACTTCCGCTCCTGGTTTAAATGAACATGGATCCAGTTTTTCGGAAAATTC  
228Lneomexicanum TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGCTCCATTTTTTCAGAAAATGT  
231Lcompactum TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGTTCCATTTTTTCAGAAAATGT  
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233Laristatum TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGCTCCATTTTTTCAGAAAATGT  
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236Lrigidumberla TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGTTCCATTTTTTCAGAAAATGT  
238Lpratense TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGTTCCATTTTTTCAGAAAATGT  
241Lrigidumberla TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGTTCCATTTTTTCAGAAAATGT  
242Lrupestre TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATAGATATGGCTCCATTTTTTCAGAAAATGT  
246LrigidumKerr TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGTTCCATTTTTTCAGAAAATGT  
253Lelongatum TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGTTCCATTTTTTCAGAAAATGT  
256Lmedium TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
257Lmedium TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
258Lquadrifolium TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
259Lheterostylum TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
266Lcarteri TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGTTCCATTTTTTCAGAAAATGT  
268Larenicola TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATAGATATGGCTCCATTTTTTCAGAAAATGT  
269Lwestii TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
270Lorizabae TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
291Lrigidumfilif TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGTTCCATTTTTTCAGAAAATGT  
293Lquatemalense TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATGT  
294Lcatharticum TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGATCCCTTTTTTCGGAAAATCT  
302Lsubteres TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATCGATATGGCTCCATTTTTTCAGAAAATGT  
307Lalatum TCGGGAGTATATTTATGTAGTTCTTATGCTCGTGGTTTAAATAGATATGGCTCCATTTTTTCAGAAAATGT  
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244Lhudsonioides TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGCTCCATTTTTTCAGAAAATGT  
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250Lhudsonioides TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGCTCCATTTTTTCAGAAAATGT  
AlLrigidumpuberu TCGGGAGTATATTTATGTAGTTACTTATGCTCGTGGTTTAAATAGATATGGTTCCATTTTTTCAGAAAATGT  
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L43Lcatharticum TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGATCCCTTTTTTCGGAAAATCT  
L57Lflavum -----  
LPB2Loligophyllu TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCAATTTTTTCAGAAAATGT  
PM1983Sdigynum TCGGGAGTATATTTATATAGTTACTTATGCTCGTGGTTTAAATGGATATGGCTCCATTTTTTCAGAAAATTT



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 091Loligophyllum ACCGAATCATTTTCATTATTTTCGGCTACTGATTCTGAACAACATATACTTTTTAGGTATAACAACATTTTGTA  
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 165Lrzedowskii -----  
 171Lrlinoides ACCGAATCATTTTCATTATTTTCGGCTACTGATTCTGAACAACATATACTTTTTAGGTATAACAACATTTTGTA  
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067Lthunbergii -----  
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067Lthunbergii -----  
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165Lrzedowskii -----  
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195Lchamissonis CACTCGCTTCTGGATAAAAAGATCCCGCTTGTTTGCATTTATTACGACTCTTTTTTCACGAGT-----ATTG  
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200Lcarneum CACTCGCTTCTGGATAAAAAGATCCCGCTTGTTTGCATTTATTACGACTCTTTTTTCACGAGTATTGGAATTG  
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232Laustrale CACTCGTTTCTGGATAAAAAGATCCCGCTTGTTTGCATTTATTACGACTCTTTTTTCACGAGTATTGGAATTG  
233Laristatum CACTCGTTTCTGGATAAAAAGATCCCGCTTGTTTGCATTTATTACGACTCTTTTTTCACGAGTATTGGAATTG  
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 067Lthunbergii -----  
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 L29Lmaritimum TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCACTTTTATCA  
 L4Lusitatisimum TATATATGAAAATGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTATTTACGATCAACCTTTTCTCG  
 L41Lviscosum -----TTTTCTCG  
 L43Lcatharticum TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCACTTTTATCA  
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 LPB2Loligophyllu TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCACTTTTATCA  
 PM1983Sdigynum TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCACTTTTATCA  
 SA1Lafricanum TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCACTTTTATCA  
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 SA11LinumAgulhas TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCACTTTTATCA  
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 SA14Lvillosum TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCACTTTTATCA  
 SA17Lthesioides TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCACTTTTATCA  
 SA19Lcomptonii TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCAC-----  
 SA3Lacuticarpum TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCACTTTTATCA  
 SA4Lgracile TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCACTTTTATCA  
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 SA8Ladustum TATATATGAAAACGAATCCATCTTCTTTTTCTTCGTAACCAATCTTTTCATTTACGATCCACTTTTATCA  
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194Lprostratum GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
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196Lerigeroides GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
200Lcarneum GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
201Lpringlei GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
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223Lkingii -----  
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302Lsubteres -----  
307Lalatum GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
310Llasiocarpum -----  
244Lhudsonioides GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
214Lbienne CTTCCTTCTTGAGCGAAATTTGTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTACTCATTTTGGGG  
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F2Lcruciata GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
F5Lnelsonii GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
F6Ltenellum GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
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SA14Lvillosum GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
SA17Lthesioides GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
SA19Lcomptonii -----  
SA3Lacuticarpum GTTTCCTTCTTGAGCGAAAGTATTTCTATGGAAAAATAGAACAATTTGCGGAAGTCTTTGCTAATTGTTTTTCG  
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165Lrzedowskii GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
171Rlinoides GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
183Lpuberulum GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATATATTCTAGCTTC  
184Lvernale GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
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193Lbrevifolium GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
194Lprostratum GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
195Lchamissonis GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
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221Lburkartii GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
223Lkingii -----  
224Lmonogynum GATCCGCTTATGGTTATTTAAAGGATACTTTTCATGCATTATGCTAGATATCAAGGAAAATTTATTTTTCAGCTTC  
228Lneomexicanum GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
231Lcompactum GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATATATTCTAGCTTC  
232Laustrale GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATATATTCTAGCTTC  
233Laristatum GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATATATTCTAGCTTC  
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294Lcatharticum GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
302Lsubteres -----  
307Lalatum GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATATATTCTAGCTTC  
310Llasiocarpum -----  
244Lhudsonioides GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
214Lbienne GATCCGCTTATGGATATTTAAAGGATACTTTTCATGCATTATGCTAGATATCAAGGAAAATTTTTCAGCTTC  
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Fl1Lbahamensecora GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
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SA10Lesterhuysen GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
SA11LinumAgulhas GACCAGCCTATGGCTGTTTAAAGGATCCGTTTCATGCATTATGCTAGATATCAAGGAAAATACATTCTAGCTTC  
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149Lflavum AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
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259Lheterostylum AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
266Lcarteri AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
268Larenicola AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
269Lwestii AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
270Lorizabae AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTGA  
291Lrigidumfilif AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
293Lquatemalense AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTGA  
294Lcatharticum AAAGGGTACGACGGCCCTTCTAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
302Lsubteres -----  
307Llalatam AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
310Llasiocarpum -----  
244Lludsonioides AAAGGGTACACCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
214Lbienne AAAGGGTACGCTCC---TTTCAATTAATAAATGGAATATTATCTTGTCAATTTATGGCAATGTCATTTTTTA  
249LinumBrackenr AAAGGGTACGCGC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
250Lludsonioides AAAGGGTACACCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
A1Lrigidumpuberu AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
F1Lbahamensecora AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
F2Lcruciata AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
F5Lnelsonii AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
F6Ltenellum AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
F7Lscabrellum AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
L17Ltenuifolium AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
L29Lmaritimum AAAGGGTACGCCCT---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTGA  
L4Lusitatissimum AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
L41Lviscosum AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
L43Lcatharticum AAAGGGTACGACGGCCCTTCTAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA  
L57Lflavum AAAGGGTACGCCCC---TTCGAATTAATAAATGGAATATTACCTTGTCAATTTATGGCAATGTCATTTTTTA



124Lfloridanum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
130Lvirginianum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
132Lsulcatum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
137Lvolskensis TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACAGTCTATTCACCTTTTATAGGCTATCT  
139Lholstii TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACAGTCTATTCACCTTTTATAGGCTATCT  
143Lorientale TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
146Lnervosum TGTGTGGTCTCAACCGAAAACATCTATTTAAATTCATTATCTAAACATTCTATCCACTTTTATAGGCTATCT  
148Lhypericifoli TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
149Lflavum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
151Lcatharticum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
156Ltenuifolium TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
157Ltauricum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
163Llegans TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTAGGCACTTTTATAGGCTATCT  
164Larboreum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
165Lrzedowskii TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
171Lrlinoides TGTGTGGTCTCGCAACCAGAAAACATTTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
183Lpuberulum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
184Lvernale TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
186Llundellii TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
187Lcremonophyllu TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
193Lbrevifolium TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
194Lprostratum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
195Lchamissonis TGTGTGGTCTCAACCAGAAAACATCTATTTAAATCCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
196Lerigeroides TGTGTGGTCTCAACCAGAAAACATCTATTTAAATCCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
200Lcarneum TGTGTGGTCTCAACCAGAAAACATCTATTTAAATCCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
201Lpringlei TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
206Cselaginoides TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
210bLmacraeimacr TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
215Lchamissonis TGTGTGGTCTCAACCAGAAAACATCTATTTAAATCCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
219Lscoparium TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
220Cselaginoides TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
221Lburkartii TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
223Lkingii -----  
224Lmonogynum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACGTATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
228Lneomexicanum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
231Lcompactum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
232Laustrale TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
233Laristatum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
234Lrigidumberla TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
236Lrigidumberla TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
238Lpratense TGTGTGGTCTCAACCGAAAACATCTATTTAAATTCATTATCCAAAGCATTCTGTCCACTTTTATAGGCTATCT  
241Lrigidumberla TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
242Lrueprestre TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
246LrigidumKerr TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
253Lelongatum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
256Lmedium TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
257Lmedium TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
258Lquadrifolium TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCGCTTTTATAGGCTATCT  
259Lheterostylum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCGCTTTTATAGGCTATCT  
266Lcarteri TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
268Larenicola TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTTATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
269Lwestii TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
270Lorizabae TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
291Lrigidumfilif TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
293Lquatemalense TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
294Lcatharticum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
302Lsubteres -----  
307Lalatum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
310Llasiocarpum -----  
244Lhudsonioides TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAAGCATTCTATCCACTTTTATAGGCTATCT  
214Lbienne TGTGTGGTCTCAACCAGAAAACATCTATTTAAACATATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
249LinumBrackenr TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
250Lhudsonioides TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAAGCATTCTATCCACTTTTATAGGCTATCT  
AlLrigidumpuberu TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
F1Lbahamensecora TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
F2Lcruciata TGTGTGGTCTCAACCAGAAAACATCTATTTAAACGATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
F5Lnelsonii TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT  
F6Ltenellum TGTGTGGTCTCAACCAGAAAACATCTATTTAAACTCATTATCCAAACATTCTATCCACTTTTATAGGCTATCT



072Ltrigynum	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
075Lstriatum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
076Lmarginale	TTCAAGTCTAGGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATAGATAATAC
083Sdigynum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
089Llittorale	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
091Lloliophyllum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
123Lkingii	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
124Lfloridanum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
130Lvirginianum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
132Lsulcatum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
137Lvolskensis	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
139Lholstii	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
143Lorientale	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
146Lnervosum	TTCAAGTCTAGGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTATCTAATAGATAATAG
148Lhypericifoli	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATAGATAATAC
149Lflavum	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
151Lcatharticum	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
156Llenuifolium	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
157Ltauricum	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
163Lpulegans	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
164Larboreum	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
165Lrzedowskii	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
171Lrlinoides	TTCAAGTCTACGATTAAATCCTTCAGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATAGATAATAC
183Lpuberulum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
184Lvernale	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
186Llundellii	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATGGATAATAC
187Lcremnophyllum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
193Lbrevifolium	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
194Lprostratum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
195Lchamissonis	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
196Lrigeroides	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
200Lcarneum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
201Lpringlei	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
206Cselaginoides	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
210bLmacraeimacr	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
215Lchamissonis	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
219Lscoparium	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
220Cselaginoides	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
221Lburkartii	TTCAAGTGTACAAATTAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
223Lkingii	-----
224Lmonogynum	TTCAAGTCTAGGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATAGATAATAC
228Lneomexicanum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
231Lcompactum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATGGATAATAC
232Laustrale	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
233Laristatum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
234Lrigidumberla	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATGGATAATAC
236Lrigidumberla	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATGGATAATAC
238Lpratense	TTCAAGTCTAGGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATAGATAATAG
241Lrigidumberla	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATGGATAATAC
242Lrupestre	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
246LrigidumKerr	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATGGATAATAC
253Lelongatum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATGGATAATAC
256Lmedium	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
257Lmedium	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
258Lquadrifolium	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
259Lheterostylum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
266Lcarteri	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATGGATAATAC
268Llarenicola	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
269Lwestii	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
270Llorizabae	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
291Lrigidumfilif	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATGGATAATAC
293Lquatemalense	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
294Lcatharticum	TTCAAGTCTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
302Lsubteres	-----
307Llatum	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
310Llasiocarpum	-----
244Lhudsonioides	TTCAAGTGTACGATTAAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
214Lbienne	TTCAAGTCTAGAAATCAATCCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTCTAATAGATAATAC

249LinumBrackenr	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
250Lhudsonioides	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
AlLrigidumpuberu	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
F1Lbahamensecora	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
F2Lcruciata	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
F5Lnelsonii	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
F6Ltenellum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
F7Lscabrellum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
L17Ltenuifolium	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
L29Lmaritimum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
L4Lusitatisimum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
L41Lviscosum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
L43Lcotharticum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
L57Lflavum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
LPB2Loligophyllu	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
PM1983sdigynum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA1Laficanum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA10Lesterhuysen	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA11LinumAgulhas	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA12Laethiopicum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA13Linumspwindb	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA14Lvillosum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA17Lthesioides	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA19Lcomptonii	-----
SA3Lacuticarpum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA4Lgracile	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA5Lgracile	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA6Lgracile	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA7Linumdehoop	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA8Ladustum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SA9Lesterhuysena	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
SD1Lrigidumcompa	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
Gyp1YesoHills	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
400Lrigidumrigid	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
162Ldecumbens	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC
401Lintercursum	TTCAAGTGTACGATTAATAACCTTCGGTCTTACGAAGTCAAATGGTAGAAAAATCTTTTATAATGGATAATAC



049Cselaginoides TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
 055Laethiopicum TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
 059Lcomptonii TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
 061Lesterhuysena TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
 066Ltenue TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
 062Lheterostylum -----  
 067Lthunbergii -----  
 072Ltrigynum TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
 075Lstriatum TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
 076Lmarginale TAAGCATAAATTTGATGCAAAATTTCCAATTTTCTTTTCGTTGATTGCAGCCTTGGAAGAACCAAAATTTGTAA  
 083Sdigynum TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
 089Llittorale TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTAATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
 091Lloliophyllum TGCGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
 123Lkingii TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
 124Lfloridanum TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAA  
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 143Lorientale TACGCAGAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGGAGCCTTGGAAGAACCAAAATTTGTAC  
 146Lnervosum AAAGCATAAATTTGATACAATAATTCCAATTTTCTTTTCGTTGATTTCAGCCTTGGAAGAACCAAAATTTGTAA  
 148Lhypericifoli TAGGCATAAATTTGATACAATAATTCCAATTCCTTTTCGTTGATTGCAGCCTTGGAAGAACCAAAATTTGTAA  
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 223Lkingii -----  
 224Lmonogynum TAAGCATAAATTTGATGCAAAATTTCCAATTTTCTTTTCGTTGATTGCAGCCTTGGAAGAACCAAAATTTGTAA  
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038Llongipes CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 040Lcruciata CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 041Laristatum CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 042Lflagellare CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 044Reinwardtiaian CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACGAGAAAAACAGA  
 047Hdrymarioides CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 048Hmicranthum CGACTGTACGCAGTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 049Cselaginoides CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 055Laethiopicum CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 059Lcomptonii CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 061Lesterhuysena CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 066Ltenue TGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 062Lheterostylum --ACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 067Lthunbergii --ACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 072Ltrigynum TG-----GG-TGGGGTCGGAATTG-TGG-AGAG-TTATTACCGACGAAAA-CAGA  
 075Lstriatum CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 076Lmarginale CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACGAGAAAAACGGA  
 083Sdigynum CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 089Llittorale CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 091Loligophyllum CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 123Lkingii CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 124Lfloridanum CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 130Lvirginianum CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 132Lsulcatum CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 137Lvolskensisii CGACTATACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 139Lholstii CGACTGTACGCA-TTTTTTGAAGAAGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 143Lorientale CGACTGTACGCATTTTTTTGAAAAGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
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 148Lhypericifoli CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACGAGAAAAACGGA  
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 164Larboareum CGACTGTACGCATTTTTTTGAAAAGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 165Lrzedowskii CGACTGTACGCA--TTTTTGAAGAAGTTGGGGTCGGGATTGTTGGAAGAATTTTTTACGAGAAAAACGGA  
 171Rlinoides TGACTGTACGCATTTTTTT-AAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGAGAAAAACAGA  
 183Lpuberulum CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 184Lvernale CGACTGTACGCATTCTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 186Llundellii CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 187Lcremnophyllu CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 193Lbrevifolium CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 194Lprostratum CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 195Lchamissonis CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 196Lerigeroides CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 200Lcarneum CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 201Lpringlei CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 206Cselaginoides CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 210bLmacraeimacr CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
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 219Lscoparium CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 220Cselaginoides CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 221Lburkartii CG-----CGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 223Lkingii --ACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 224Lmonogynum CG-----GTTGGGGTCGGAATTGTTGGAAGAATTTTTTACGAGAAAAACGGA  
 228Lneomexicanum CG-----GGTGGGGTCGGAATTGTTGGAAGAATTTTTTACGAGAAAAACGGA  
 231Lcompactum CG-CTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 232Laustale CGACTGTACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 233Laristatum CGACTGTACGCATTTTTTTG- AAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 234Lrigidumberla CG-----AAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 236Lrigidumberla CG---TACGCATTTTTTTGAAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 238Lpratense CGACTGTACGCATTTTTTT-AAAAGGTTGGGGTCGGGATTGTTGGAAGAATTTTTTACGAGAAAAACGGA  
 241Lrigidumberla CG---TACGCATTTTTTTG- AAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 242Lrupestre CGACTGTACGCATTTTTTT-AAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 246LrigidumKerr CG-----TTTTTTT- AAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 253Lelongatum CGACTGTACGCATTTTTTT-AAAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 256Lmedium CGACTGTACGCATTTTTTT- AAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA  
 257Lmedium CGACTGTACGCATTTTTTT- AAAGGTTGGGGTCGGAATTGTTGGAAGAATTTTTTACCGACGAAAAACAGA



027Lberlandierib TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
028Lberlandierif TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
032Lhudsonioides TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
034Lrupestre TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGAGTTTGGTATTTGGATA  
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038Llongipes TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
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041Laristatum TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
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049Cselaginoides -----  
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089Llittorale -----  
091Loligophyllum -----  
123Lkingii TTCTTTCTTTGATTCCAAAAACCTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
124Lfloridanum TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
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143Lorientale TTCTTTCTTT-----AACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
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148Lhypericifoli TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
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156Ltenuifolium TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
157Ltauricum TTCTTTCTTT-----AACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
163Llelegans TTCTTTCTTT-----AACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
164Larboreum TTCTTTCTTT-----AACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
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171Rlinoides TTCTTTCTTTGATTCCAACAACCTTCAGCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
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186Llundellii TTCTTTCTTTGAGTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
187Lcremnophyllu -----  
193Lbrevifolium -----  
194Lprostratum -----  
195Lchamissonis -----  
196Lerigeroides -----  
200Lcarneum -----  
201Lpringlei TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGAGTTTGGTATTTGGATA  
206Cselaginoides -----  
210bLmacraeimacr -----  
215Lchamissonis -----  
219Lscoparium -----  
220Cselaginoides -----  
221Lburkartii -----  
223Lkingii TTCTTTCTTTGATTCCAAAAACCTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
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231Lcompactum TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
232Laustale TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
233Laristatum TTCTTTCTTTGATTCCAAGAACTTCGCTCTATTTCCAAAAGATTATATAAAGGACGGGTTTGGTATTTGGATA  
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009Lschiedeanum -----  
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011Lsuffruticosu TTGTTTGTATCAATGAACTGGTCCATGATAAAAAATTTGTTTATCAAACATGGAAATAGAAATTTTCCATAAA  
012Lstrictum TTGTTTGTATCAATGAACTGGTCCATGATAAAAAATTTGTTTATCAAACATGGAAATAGAAATTTTCCATAAA  
013Lnarbonense TTGTTTGTATCAATGAACTGGCCCATGATAAAAAATTTTATGAAACATGGAAATCGAAATTTCCATAAA  
022Hclevelandii TTGTTTGTATCAATGAACTGGCCCATGATAAAAAATTTCTTTATCAAACATGGAAATGAAATTTTCCATAAA  
025Hdisjunctum TTGTTTGTATCAATGAACTGGCCCATGATAAAAAATTTCTTTATCAAACATGGAAATGAAATTTTCCATAAA  
027Lberlandierib TTGTTTGTATCAATGAACTGGCCCATGATCAAAATTTCTTTATCAAACATGGAAATAGAAATTTTCCATAAA  
028Lberlandierif TTGTTTGTATCAATGAACTGGCCCATGATCAAAATTTCTTTATCAAACATGGAAATAGAAATTTTCCATAAA  
032Lhudsonioides TTGTTTGTATCAACGAACTGGCCCATGATCAAAATTTCTTTATCAAACATGGAAATAGAAATTTTCCATAAA  
034Lrueprestre TTGTTTGTATCAATGAACTGGCCCATGATCAAAATTTCTTTATCAAACATGGAAATAGAAATTTTCCATAAA  
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089Llittorale -----  
091Loligophyllum -----  
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187Lcremnohyllum -----  
193Lbrevifolium -----  
194Lprostratum -----  
195Lchamissonis -----  
196Lerigeroides -----  
200Lcarneum -----  
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210bLmacraeimacr -----  
215Lchamissonis -----  
219Lscoparium -----  
220Cselaginoides -----  
221Lburkartii -----  
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244Lhudsonioides -----  
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206Cselaginoides -----  
210bLmacraeimacr -----  
215Lchamissonis -----  
219Lscoparium -----  
220Cselaginoides -----  
221Lburkartii -----  
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186Llundellii	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
187Lcremnophyllu	-----
193Lbrevifolium	-----
194Lprostratum	-----
195Lchamissonis	-----
196Lerigeroides	-----
200Lcarneum	-----
201Lpringlei	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTAGGGGAAACTGAGCTTT
206Cselaginoides	-----
210bLmacraeimacr	-----
215Lchamissonis	-----
219Lscoparium	-----
220Cselaginoides	-----
221Lburkartii	-----
223Lkingii	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTAGAGGAAACTGAGCTTT
224Lmonogynum	ATTGAGCAACCAAGTATTCAACTTTCTTCTGAGGTATGGGGTGAGTTCTGTCTAGGGGAAACTGAGCTTT
228Lneomexicanum	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-AGGAAACTGAGCTTT
231Lcompactum	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGACCTTT
232Laustrale	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
233Laristatum	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTAGGGGAAACTGAGCTTT
234Lrigidumberla	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGACCTTT
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238Lpratense	ATTGAGCAACCAAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
241Lrigidumberla	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGACCTTT
242Lrupestre	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
246LrigidumKerr	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGACCTTT
253Lelongatum	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGACCTTT
256Lmedium	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
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259Lheterostylum	ATTGAGCAACTGAGTATTCAACTTTCTTCTAAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
266Lcarteri	-----
268Larenicola	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
269Lwestii	ATTCAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
270Lorizabae	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
291Lrigidumfilif	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTAGGGGAAACTGACCTTT
293Lquatemalense	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
294Lcatharticum	ATTGAGCAACGAGTATTCCACTTTCTTCTGAGGTATGGGT-----T-CCGTCTA-GGGAAACTGAGCTTT
302Lsubteres	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTAGGGGAAACTGAGCTTT
307Llalatium	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTAGGGGAAACTGAGCTTT
310Llasiocarpum	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTAGGGGAAACTGAGCTTT
244Lhudsonioides	-----
214Lbienne	ATTGAGCAACCAAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
249LlinumBrackenr	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
250Lhudsonioides	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
Al1rigidumpuberu	ATTGAGCAATTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGACCTTT
F1Lbahamensecora	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
F2Lcruciata	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
F5Lnelsonii	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
F6Ltenellum	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
F7Lscabrellum	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTAGGGGAAACTGAGCTTT
L17Ltnuifolium	ATTGAGCGACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCCGTCTA-GGGAAACTGAGCTTT
L29Lmaritimum	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTACGGGG-----TTCCGTCTA-GGGAAACTGAGCTTT
L4Lusitatissimum	ATTGAGCAACCAAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
L41Lviscosum	ATTGAGCAACCAAGTATTCAACTTTCTTCTGAGGTATGGGG-----CTCCGTCTA-GGGAAACGAGCTTT
L43Lcatharticum	ATTGAGCAACGAGTATTCCACTTTCTTCTGAGGTATGGG-----TTCCGTCTA-GGGAAACTGAGCTTT
L57Lflavum	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCCGTCTA-GGGAAACGAGCTTT
LPB2Loligophyllu	-----
PM1983Sdigynum	ATTGAGCAACTGAGTATTCAACTTTCTTCTGAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
SA1Lafricanum	ATTGAGCAACTGAGTATTCAACTTTCTTCTAAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
SA10Lesterhuysen	ATTGAGCAACTGAGTATTCAACTTTCTTCTAAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
SA11LinumAgulhas	ATTGAGCAACTGAGTATTCAACTTTCTTCTAAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
SA12Laethiopicum	-----
SA13LinumspWindb	ATTGAGCAACTGAGTATTCAACTTTCTTCTAAGGTATGGGG-----TTCTGTCTAGGGGAAACTGAGCTTT
SA14Lvillosum	ATTGAGCAACTGAGTATTCAACTTTCTTCTAAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
SA17Lthesioides	ATTGAGCAACTGAGTATTCAACTTTCTTCTAAGGTATGGGG-----TTCTGTCTA-GGGAAACTGAGCTTT
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193Lbrevifolium -----ATACCTTACATAATGAATCTGATA  
194Lprostratum -----ATACCTTACATAATGAATCTGATA  
195Lchamissonis -----ATACCTTACATAATGAATCTGATA  
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LPB2Loligophyllu -----ATACCTTACATAATGAATCTGATA  
PM1983Sdigynum AGTTGTATACATAGGAAAAGCCGTGTGCAATGAAAAATGCAAGCACGGATACCTTACATAATGAATCTGATA  
SA1Lafricanum AGTTGTATACATAGGAAAAGCCGTGTGCAATGAAAAATGCAAGCACGGATACCTTACATAATGAATCTGATA



132Lsulcatum GCTTCGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT  
137Lvolkensis GCTTTGATAGGTATACTAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCTCAAAAAGATATTAAGAGAAGT  
139Lholstii GCTTTGATAGGTATACTAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCTCAAAAAGATATTAAGAGAAGT  
143Lorientale GCTTTGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCTCAAAAAGATATTAAGAGAAGT  
146Lnervosum GCTTTGGTAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCTCTTGCTCAAAAAGATATTAAGAGAAGT  
148Lhypericifoli GCTTTAGTAGGTATAATAACAGTACTTTTAGGGGCTACTTTAGCTCTTGCTCAAAAAGATATTAAGAGAAGT  
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157Ltauricum GCTTTGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCTCAAAAAGATATTAAGAGAAGT  
163Llegans GCTTTGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCTCAAAAAGATATTAAGAGAAGT  
164Larboreum GCTTTGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCTCAAAAAGATATTAAGAGAAGT  
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200Lcarneum GCTTTGATAGGTATAATAACAGTCTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT  
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259Lheterostylum -----  
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310Llasiocarpum GCTTCGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT  
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214Lbienne GCTTTGGTAGGAATAATAACCGTACTTTTAGGAGCTACTTTAGCTCTTGCTCAAAAAGATATTAAGAGAAGT  
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L29Lmaritimum	GCTTTGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCTCAAAAAGATATTAAGAGAAGT
L4Lusitissimum	GCTTTGGTAGGAATAATAACCGTACTTTTAGGAGCTACTTTAGCTCTTGCTCAAAAAGATATTAAGAAAAAGT
L41Lviscosum	GCTTTGATAGGTATAATAACAGTACTTTTAGGGCTACTTTAGCTCTTGCTCAAAAAGATATTAAGAGAAGT
L43Lcoatharticum	GCTTTGTAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCTCAAAAAGATATTAAGAGAAGT
L57Lflavum	GCTTTGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCTCAAAAAGATATTAAGAGAAGT
LPB2Loligophyllu	GCTTTGTAGGGATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT
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SA1Laficanum	GCTTTGTAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT
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SA19Lcomptonii	GCTTTGTAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT
SA3Lacuticarpum	GCTTTGTAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT
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SA6Lgracile	GCTTTGTAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT
SA7Linumdehoop	GCTTTGTAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT
SA8Ladustum	GCTTTGTAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT
SA9Lesterhuysena	GCTTTGTAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT
SD1Lrigidumcompa	GCTTCGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAGAGATATTAAGAGAAGT
Gyp1YesoHills	GCTTCGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT
400Lrigidumrigid	GCTTCGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAGAGATATTAAGAGAAGT
162Ldecumbens	GCTTYGATAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCTCAAAAAGATATTAAGAGAAGT
401Lintercursum	GCTTTGTAGGTATAATAACAGTACTTTTAGGAGCTACTTTAGCGCTTGCGCAAAAAGATATTAAGAGAAGT

076Lmarginale	TTAGCCTATTCTACAATGTCTCAATTGGGTTATATGATGTTAGCTTTAGGAATGGGATCGTATCGGGCAGCG
083Sdigynum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGCTCGTATCGAGCGGCC
089Llittorale	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
091Loligophyllum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
123Lkingii	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
124Lfloridanum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
130Lvirginianum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
132Lsulcatum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
137Lvolskensis	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
139Lholstii	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
143Lorientale	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
146Lnervosum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGGGTGCG
148Lhypericifoli	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
149Lflavum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
151Lcatharticum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
156Ltenuifolium	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
157Ltauricum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
163Lelegans	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
164Larboreum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
165Lrzedowskii	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
171Rlinoides	TTAGCCTATTCTACAATGTCTCAATTGGGTTATATGATGTTAGCTTTAGGTATGGGATCTTACCGCGCGGCC
183Lpuberulum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
184Lvernale	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
186Llundellii	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
187Lcremophyllum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
193Lbrevifolium	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
194Lprostratum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
195Lchamissonis	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
196Lerigeroides	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
200Lcarneum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
201Lpringlei	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
206Cselaginoides	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
210bLmacraeimacr	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
215Lchamissonis	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
219Lscoparium	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
220Cselaginoides	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
221Lburkartii	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
223Lkingii	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
224Lmonogynum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATATGATGTTAGCTTTAGGAATGGGATCGTATCGGCGAGCG
228Lneomexicanum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
231Lcompactum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
232Laustale	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
233Laristatum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
234Lrigidumberla	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
236Lrigidumberla	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
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241Lrigidumberla	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
242Lrupestre	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
246LrigidumKerr	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
253Lelongatum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
256Lmedium	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
257Lmedium	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
258Lquadrifolium	-----
259Lheterostylum	-----
266Lcarteri	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
268Larenicola	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
269Lwestii	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
270Lorizabae	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
291Lrigidumfilif	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
293Lquatemalense	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
294Lcatharticum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
302Lsubteres	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
307Lalatum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
310Llasiocarpum	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
244Lhudsonioides	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
214Lbienne	TTAGCCTATTCTACAATGTCTCAATTGGGTTATATGATGTTAGCTTTAGGTATGGGATCTTATCGGCGAGCG
249LinumBrackenr	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC
250Lhudsonioides	TTAGCCTATTCTACAATGTCTCAATTGGGTTATACGATGTTAGCTTTAGGTATGGGATCTTATCGAGCGGCC



0055Laethiopicum CTTTTTCATTTGATTACTCATGCGCTATTCGAAAGCCTTATTGTTTTTAGGATCTGGATCCATTATTCATTCA  
0059Lcomptonii CTTTTTCATTTGATTACTCATGCGCTATTCGAAAGCCTTATTGTTTTTAGGATCTGGATCCATTATTCATTCA  
0061Lesterhuysena CTTTTTCATTTGATTACTCATGCGCTATTCGAAAGCCTTATTGTTTTTAGGATCTGGATCCATTATTCATTCA  
0066Ltenue CTTTTTCATTTGATTACTCATGCGCTATTCGAAAGCCTTATTGTTTTTAGGATCTGGATCTATTATTCATTCA  
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256Lmedium CTTTTTCATTTGATTACTCACGCGCTATTCGAAAGCCTTATTGTTTTTAGGATCTGGATCCATTATTCATTCA  
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258Lquadrifolium -----  
259Lheterostylum -----  
266Lcarteri CTTTTTCATTTGATTACTCATGCGCTATTCGAAAGCCTTATTGTTTTTAGGATCTGGATCCATTATTCATTCA  
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 041Laristatum ATGGAAGTATTGTTGGCTATTCTCCAGATAAGAGTCAAAATATGGTTCCTATGGGTGGTTACGAAAACAC  
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 256Lmedium ATGGAAGTATTGTTGGCTATTCTCCAGATAAGAGTCAAAATATGGTTCCTATGGGTGGTTAAGAAAACAC  
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 258Lquadrifolium -----



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 032Lhudsonioides GTGCCAATTACAAAACTACTTTTTATTAGGAACCTTTTCACTTTGTGGTGTTCGCCCCCTCGCCTGCTTT  
 034Lrupestre GTGCCAATTACAAAACTGCTTTTTATTAGGAACCTTTTCACTTTGTGGTGTTCGCCCCCTCGCCTGCTTT  
 035Lrigidumrigid GTGCCAATTACAAAACTACTTTTTATTAGGAACCTTTTCACTTTGTGGTGTTCGCCCCCTCGCCTGCTTT  
 036Lrupestre GTGCCAATTACAAAACTGCTTTTTATTAGGAACCTTTTCACTTTGTGGTGTTCGCCCCCTCGCCTGCTTT  
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401Lintercursum TGGTCCAAAGACGAAATTCCTTAATGCTAGTTGGTTATATTACCTATTTTT-----

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	000
Taxon/Node	111222222222333333333344444444445555555555666666666677777777778888888888 789012345678901234567890123456789012345678901234567890123456789012345678901234567890123456789012345678
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010Lewisii	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
011Lsuffruticosu	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTGCCTAAAGACAGAATCAAA-CAGGATAGGTGC
012Lstrictum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
013Lnarbonense	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
022Hclevelandii	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
025Hdisjunctum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
027Lberlandierib	-----
028Lherlandierif	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGCCAAAAATAAAA-CAGGATAGGTGCA
032Lhudsonioides	-----
034Lrupestre	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
035Lrigidumrigid	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGCCAAAAATAAAA-CAGGATAGGTGCA
036Lrupestre	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
037Mexicanum	C-AAAGGGGCAATCCTGAGCCAAATCC-TG-----CTCCTTAAAGACAGAATAAAA-CAGGATAGGTGCA
038Llongipes	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
040Lcruciata	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
041Laristatum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGCCAGAATAAAA-CAGGATAGGTGCA
042Lflagellare	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
044Reinwardtiaia	--AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCATTCTAAGACAGAATAAAA-CAGGATAGGTGCA
047Hdymarioides	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
048Mmicranthum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
049Cselinaginoides	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
055Laethiopium	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
059Lcomptonii	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
061Lesterhuysena	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
066Ltenue	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
062Lheterostylum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
067Lthunbergii	C-AAAGGGGTAATCCTGAGCCAAATCC-TGTTTT-CCG-----
072Ltrigynum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
075Lstriatum	-----TTCTTAAAGACAGAATAAAA-CAGGATAGGTGCA
076Lmarginale	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAGACAGACTAAAA-CAGGATAGGTGCA
083Sdigynum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-C--TTCTTAAAGACAGAATAAAA-CAGGATAGGTGCA
089Llittorale	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
091Loligophyllum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
123Lkingii	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
124Lfloridanum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
130Lvirginianum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
132Lsulcatum	-----
137Lvolskensi	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
139Lholstii	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
143Lorientale	T-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
146Lnervosum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-ACGTTTCAGAAAGACAGACTAAAA-CAGGATAGGTGCA
148Lhypericifoli	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-TCGTT-----AAAGATAAAA-CAGGATAGGTGCA
149Lflavum	T-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
151Lcatharticum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
156Ltenuifolium	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTGCCTAAAGACAGAATCAA-CAGGATAGGTGCA
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163Lelegans	T-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
164Larboreum	TGAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
165Lrzedowskii	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCAGAAAGACAGACTAAAA-CAGGATAGGTGCA
171RLnoides	-----
183Lpuberulum	-----GGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGCCAGAATAAAA-CAGGATAGGTGCA
184Lvernale	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGCCAGAATAAAA-CAGGATAGGTGCA
186Lludelli	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGCCAAAAATAAAA-CAGGATAGGTGCA
187Lcremnohyllu	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
193Lbrevifolium	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
194Lprostratum	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
195Lchamissonis	C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTTCCTAAAGACAGAATAAAA-CAGGATAGGTGCA
196Lerigeroides	-----

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 201Lpringlei C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
 206Cselaginoides C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
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 231Lcompactum C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGCCAAAATAAAA-CAGGATAGGTGCA  
 232Laustrale -----TTCTAAAGCCAGAATAAAA-CAGGATAGGTGCA  
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 302Lsubteres C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
 307Lalatum -----GGATAGGTGCA  
 310Llasiocarpum -----  
 244Lhudsonioides -----  
 214Lbienne C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGACTAAAA-CAGGATAGGTGCA  
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 250Lhudsonioides C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
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 L43Lcatharticum C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
 L57Lflavum T-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
 LPB2Loligophyllu C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
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 SA11LinumAgulhas C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
 SA12Laethiopicum C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
 SA13LinumspWindb C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
 SA14Lvillosum C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
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 SA19Lcomptonii C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA  
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 SA9Lesterhuysena C-AAAGGGGCAATCCTGAGCCAAATCC-TGTTTT-CCGTTCTAAAGACAGAATAAAA-CAGGATAGGTGCA



184Lvernale GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
186Llundellii GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
187Lcremophyllu GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
193Lbrevifolium GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
194Lprostratum GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
195Lchamissonis GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
196Lerigeroides -----  
200Lcarneum GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
201Lpringlei GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
206Cselaginoides GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
210bLmacraeimacr GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
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231Lcompactum GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
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233Laristatum GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
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310Llasiocarpum -----  
244Lhudsonioides -----  
214Lbienne GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
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250Lhudsonioides GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
Al1Lrigidumpuberu GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
F1Lbahamensecora GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
F2Lcruciata GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
F5Lnelsonii GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
F6Ltenellum GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
F7Lscabrellum GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
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SA1Lafricanum GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
SA10Lesterhuyzen GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
SA11LinumAgulhas GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
SA12Laethiopicum GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
SA13LinumspWindb GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
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SA17Lthesioides GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG  
SA19Lcomptonii GAGACTCGATGG-AAGCTGTTCTAACAAATGGAGGTGACCGCCTTGCGTTGGTAAAGTAAAGGAATCCTTCG





151Lcatharticum ATCAAAA-----CTCT-----  
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310Llasiocarpum -----  
244Lhudsonioides -----  
214Lbienne ATCAAAACAAAACCTTTTAACCCGATGCATAGAA-T-----CCATATGTATGGGTACTGAAATACTATATCAA  
249LinumBrackenr ATCAAAACAAAACCTGTTAACCCG-----CTCAA  
250Lhudsonioides ATCAAAAC-AAAACCTCTTAACCCG-----CTCAA  
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L41Lviscosum A-----  
L43Lcatharticum ATCAAAA-----CTCT-----  
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148Lhypericifoli -----TTCAAAATTGAAGAAAGAACC GAAT  
149Lflavum ATGATTAATGATAA----CCCGAAT-----TGAAGAAAGAATCGAAT  
151Lcatharticum -----AAAGAAAGAATCGAAT  
156Ltenuifolium ATGATTAATGATAA----CCCGAACCCCTTAAAAAGTTTTGG-----  
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244Lhudsonioides -----  
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L41Lviscosum	-----TTCAAATTTGAAGAAAGAACCGAAT
L43Lcatharticum	-----AAAGAAAGAATCGAAT
L57Lflavum	ATGATTAATGATAA---CCCGAAT-----TGAAGAAAGAATCGAAT
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PM1983Sdigynum	ATGATTAATGATACTAACCCGAATCCTTTAAAGAGTTTTGGATCGATTCCAAATTGAAGAAAGAACCGAAT
SA1Lfricanum	ATGATTAATGATACTAACCCGAATCCTTTAAAGAGTTTTGGATCGATTCCAAATTGAAGAAAGAATCGAAT
SA10Lesterhuysen	ATGATTAATGATACTAACCCGAATCCTTCAAAGAGTTTTGGATCGATTCCAAATTGAAGAAAGAATCGAAT
SA11LinumAgulhas	ATGATTAATGATACTAACCCGAATCCTTCAAAGAGTTTTGGATCGATTCCAAATTGAAGAAAGAATCGAAT
SA12Laethiopicum	ATGATTAATGATACTAACCCGAATCCTTCAAAGAGTTTTGGATCGATTCCAAATTGAAGAAAGAATCGAAT
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SA8Ladustum	ATGATTAATGATACTAACCCGAATCCTTAAAAGAGTTTTGGATCGATTCCAAATTGAAGAAAGAATCGAAT
SA9Lesterhuysena	ATGATTAATGATACTAACCCGAATCCTTCAAAGAGTTTTGGATCGATTCCAAATTGAAGAAAGAATCGAAT
SD1Lrigidumcompa	ATGATTAATGATACTAACCCGAATCCTTTAAAGAGTTTTGGATCGATTCCAAATTGAAGAAAGAATCGAAT
Gyp1YesoHills	ATGATTAATGATACTAACCCGAATCCTTTAAAGAGTTTTGGATCGATTCCAAATTGAAGAAAGAATCGAAT
400Lrigidumrigid	ATGATTAATGATACTAACCCGAATCCTTTAAAGAGTTTTGGATCGATTCCAAATTGAAGAAAGAATCGAAT
162Ldecumbens	ATGATTAATGATAA---CCCGAATCCGTCAAAGGTTTTGAATCGATTCCAAATTGAAGAAAGAACCATAAT
401Lintercursum	-----

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 156Ltenuifolium -----TCGGACGAGAATAAAGATAGAGT  
 157Ltauricum ATTAATTGATCAAATAATTCACTCCAAAGTCTGATAGATAACGGATTAATCGGACGAGAATAAAGATAGAGT  
 163Lelegans ATTAATTGATCAAATAATTCACTCCAAAGTCTGATAGATAACGGATTAATCGGACGAGAATAAAGATAGAGT  
 164Larboreum ATTAATTGATCAAATAATTCACTCCAAAGTCTGATAGATAACGGATTAATCGGACGAGAATAAAGATAGAGT  
 165Lrzedowskii TTGAATTGATCAAATAATTAACCTCCAGAGTCTGATAGATAACGGATTAATTGGACGAGAATAAAGATAGAGT  
 171Rlinoides -----  
 183Lpuberulum ATTAATTGATCAAATAATTCACTCCAAAGTCTGATAGATAACAGATTAATCGGACGAGAATAAAGATAGAGT  
 184Lvernale ATTAATTGATCAAATAATTCACTCCAAAGTCTGATAGATAACAGATTAATCGGACGAGAATAAAGATAGAGT  
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 196Lerigeroides -----  
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 228Lneomexicanum ATTAATTGATCAAATAATTCACTCCAAAGTCTGATAGATAACAGATTAATCGGACGAGAATAAAGATAGAGT  
 231Lcompactum ATTAATTGATCAAATAATTCACTCCAAAGTCTGATAGATAACAGATTAATCGGACGAGAATAAAGATAGAGT  
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 310Llasiocarpum -----  
 244Lhudsonioides -----  
 214Lbienne ATGAATTGATCAAAAATTGACTCCAAAGTCTGATAGATAACGGATTAATTGGACGAGAACAAGATAGAGT  
 249LinumBrackenr ATTAATTGATCAAATAATTCACTCCAAAGTCTGATAGATAACAGATTAATCGGACGAGAATAAAGATAGAGT  
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059Lcomptonii CCTATTCTACA-----TGTC AATATTGACAACAAGGAAATTTATAGTAAGAGGAAAATCATCCATTCTCT  
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 132Lsulcatum -----AAA--TCATCCTATTCTCT  
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 294Lcatharticum CCTATTCTACA-----TGTC AGTATTGACAACAAGGAAATTTATAGTAAGAGGAAAATCATCCTATTCTCT





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048Hmicranthum CTTTTCGTTAACGGTTTCAAATTCCTTATCTTCTCAT-TCATTGAATTCCTTTATAAAAGTATTTGGGGGT  
049Cselaginoides CTTTTCGTTAACGGTTTCAAATTCCTTATCTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG  
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196Lerigeroides -----  
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201Lpringlei CTTTTCGTTAACGGTTTCAAATTCCTTATCTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG  
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266Lcarteri CTTTTCGTTAACGGTTTCAAATTCCTTAGCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG

268Larenicola	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
269Lwestii	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
270Lorizidabae	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
291Lorizidumfilif	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
293Lquatemalense	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
294Lcatharticum	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
302Lsubteres	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
307Lalatum	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
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244Lhudsonioides	-----
214Lbienne	CTTTTCGTTAACGGTTTCAA-TCCTTATCTTTCTCAT-TCAAAGAATTCCTTTACAAAATTCCTTTGGAGTG
249LinumBrackenr	CTTTTCGTTAACGGTTTCAAATTCCTTATTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
250Lhudsonioides	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
AlRigidumpuberu	CTTTTCGTTAACGGTTTCAAATTCCTTAGCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
F1Lbahamensecora	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
F2Lcruciata	-----
F5Lnelsonii	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
F6Ltenellum	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTT--CAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
F7Lscabrellum	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
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L41Lviscosum	CTTTTCGTTAACGGATTTAAATTCCTTATCTTTCTCAT-TCATTGAATTTATTTACAAAAGTATT-GGGGTG
443Lcatharticum	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTGACAAAAGTATTTGGGGTG
L57Lflavum	CTTTTCGTTAACGGTTTAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
LPB2Loligophyllu	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
PM1983Sdigynum	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
SA1Lafricanum	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
SA10Lesterhuysen	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
SA11LinumAgulhas	CTTTTCGTTAACGATTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
SA12Laethiopicum	CTTTTCGTTAACGATTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
SA13LinumspWindb	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TTATTGAATTCCTTTACAAAAGTATTTGGGGTG
SA14Lvillosum	-----
SA17Lthesioides	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
SA19Lcomptonii	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
SA3Lacuticarpum	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
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SA6Lgracile	CTTTTCGTTAACGGTTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
SA7Linumdehoop	CTTTTCGTTAACGATTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
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SA9Lesterhuysena	CTTTTCGTTAACGATTTCAAATTCCTTATCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
SD1Lrigidumcompa	CTTTTCGTTAACGGTTTCAAATTCCTTAGCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
GyplYesoHills	CTTTTCGTTAACGGTTTCAAATTCCTTAGCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
400Lrigidumrigid	CTTTTCGTTAACGGTTTCAAATTCCTTAGCTTTCTCAT-TCATTGAATTCCTTTACAAAAGTATTTGGGGTG
162Ldecumbens	CTTTTCGTTAACGGTTTAAATTCCTTATCTTTCTTAT-TCAAAGAATTCCTTTACAAAATTCCTTTGGGGTG
401Lintercursum	-----

Taxon/Node	12345678901234567890123456789012345678901234567890123456789012
001Lstriatum	AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAAAGAACATCTTTTATTTAGCAAAACTCCCC
005Limbricatum	AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAAAGAACATCTTTTATTTAGCAAAACTCCCC
006Lbahamense	AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAAAGAACATCTTTTATTTAGCAAAACTCCCC
007Llbahamense	AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAAAGAACATCTTTTATTTAGCAAAACTCCCC
009LSchiedeanum	AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAAAGAACATCTTTTATTTAGCAAAACTCCCC
010LLewisii	AATCCAAGTTAAGTCTTGGAATAGATATAATAGACATAGAAAAGAACATGTCTTTTTAACAAAACTCCCTC
011LSuffruticosu	CACACAAGTTAAGTCTTCCTAATAGATATGATACACATAGAAAAGAACATCTTTTATTTAGCAAAACTCCCC
012Lstrictum	CACACAAGTTAAGTCTTCTTAATAGATATGATACACATAGAAAAGAACATCTTTTATTTAGCAAAACTCCCC
013Lnarbonense	AATACAAGTTGAGTCTTGTAATAGAGATAATACACATAGAAAAGAACATCTTTTATTTAGCAAAACTCCCC
022Hclevelandii	AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAAAGAACATCTTTTATTTAGCAAAACTCCCC
025Hdisjunctum	AACACAAGTTAAGTCTGGTAATAGATATGATACACATAGAAAAGAACATCTTTTATTTAGCAAAACTCCCC
027Lberlandierib	--
028Lberlandierif	AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAAAGAACATCTTTTATTTAGCAAAACTCCCC
032LBhudsonioides	--

034Lrupestre AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
035Lrigidumrigid AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCAC  
036Lrupestre AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
037Lmexicanum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
038Llongipes AA-----  
040Lcruciata AA-----  
041Laristatum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
042Lflagellare AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
044Reinwardtiain AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
047Hdrymarioides AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
048Hmicranthum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
049Cselaginoides AACACAAGTTAAGTCTTGTAATAGATATGATACATATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
055Laethiopicum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
059Lcomptonii AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
061Lesterhuysena AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
066Ltenue AACACAAGTTAAGTCTTGTAATAGATATGATATACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
062Lheterostylum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
067Lthunbergii AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
072Ltrigynum AACACAAGTTAAGTCTTGTAATAGATATGATATACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
075Lstriatum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
076Lmarginale AGTACAAGTTAAGTCTTGGAATAGAGATAATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
083Sdigynum CACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
089Llittorale AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
091Loligophyllum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
123Lkingii AAGACAAGTTAAGTCTTGTAATAGATATGATACACATTGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
124Lfloridanum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
130Lvirginianum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
132Lsulcatum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
137Lvolskensis AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
139Lholstii AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
143Lorientale AACACAAGTTAAGTCTTGTAATAGATATGATATACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
146Lnervosum AATACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATTTACATCTTAGTTTAAACAAAACCTCCCCC  
148Lhypericifoli AACAGAAGTTAAGTCTTGGAATAGATATAATATACATAGAAATGAATATCGTTTTTAAACAAAACCTCCCCC  
149Lflavum AACACAAGTTAAGTCTTGTAATAGATATGATATACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
151Lcatharticum CACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
156Ltenuifolium CACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
157Ltauricum AACACAAGTTAAGTCTTGTAATAGATATGATATACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
163Lelegans AACACAAGTTAAGTCTTGTAATAGATATGATATACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
164Larboreum AACACAAGTTAAGTCTTGTAATAGATATGATATACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
165Lrzedowskii AATCCAAGTTAAGTCTTGGAATAGATATAATAGACATAGAAATGAACATGCTTTTTTAAACAAAACCTCCCCC  
171Lrinoides CACCCAAGTTAAGTCTTGGAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
183Lpuberulum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
184Lvernale AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
186Llundellii AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
187Lcremophyllum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
193Lbrevifolium AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
194Lprostratum AACACAAGTTAAGCTTGGAATAAATAGAAACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
195Lchamissonis AA-----  
196Lerigeroides -----  
200Lcarneum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAATCCCCC  
201Lpringlei AACACAAGTTAAGTCTTGTAATAGATATAATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
206Cselaginoides AACACAAGTTAAGTCTTGTAATAGATATGATACATATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
210bLmacraeimacr AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
215Lchamissonis AA-----  
219Lscoparium AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
220Cselaginoides AACACAAGTTAAGTCTTGTAATAGATATGATACATATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
221Lburkartii AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
223Lkingii AAGACAAGTTAAGTCTTGTAATAGATATGATACACATTGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
224Lmonogynum AGTACAAGTTAAGTCTTGGAATAGAGATAATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
228Lneomexicanum AAGACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
231Lcompactum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
232Laustrale AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
233Laristatum AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
234Lrigidumberla AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
236Lrigidumberla AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
238Lpratense AATCCAAGTTAAGTCTTGGAATAGATATAGATAGACATAGAAATGAACATGCTTTTTTAAACAAAACCTCCCCC  
241Lrigidumberla AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTAGCAAAACTCCCCC  
242Lrupestre AACACAAGTTAAGTCTTGTAATAGATATGATACACATAGAAATGAACATCTTTTATTTA-----



012Lstrictum TGGAACTTGTAAATGATTAACAATCAAAAGAATTATTTCGGACTGAAAACATAAATTTAGGATATAAAACGTTTC

013Lnarbonense TGGAACTTGTAAAGGATTAACAATCCGAATCATTAT-----

022Hclevelandii TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

025Hdisjunctum RGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

027Lberlandierib -----

028Lberlandierif TGGAAATGTAAATGATTAACAATCAAAATCATTATT-----C

032Lhudsonioides -----

034Lrupestre TGGAAAGTGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

035Lrigidumrigid TGGAAATGTAAATGATTAACAATCAAAA-----CAAA-----ATTATTC

036Lrupestre TGGAAAGTGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

037Lmexicanum TGGAAATGTAAATGATTAACAATCCGAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

038Llongipes -----AGGATATAAAACGTTTC

040Lcruciata -----AGGATATAAAACGTTTC

041Laristatum TGGAAATTGGAAATGATTAACAATCAAAATCATTAT-----TC

042Lflagellare TGGAAAGTGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

044Reinwardtiain TGGAAATGTAAATGATTAACAATCCAAATCATTATTCGGACTGAAA-CTTCCAAAAGGATACAAAACGTTTC

047Hdrymarioides TGGAAATGTAAATGATTAACAATCAAAATAATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

048Hmicranthum TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

049Cselaginoides TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

055Laethiopicum TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

059Lcomptonii TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

061Llesterhuysena TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

066Ltenue CGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTAAAAAGGATATAAAACGTTTC

062Ltherostylum TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

067Lthunbergii TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

072Ltrigynum CGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTAAAAAGGATATAAAACGTTTC

075Lstriatum TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

076Lmarginale TGGAACTTGTAAAGGATTAACAATCCGAACCATTTAT-----

083Sdigynum TGGAAATGTAAATGATTAACAATCCAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

089Llittorale TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

091Loligophyllum TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

123Lkingii TGGAAATTTTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

124Lfloridanum TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

130Lvirginianum TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

132Lsulcatum TGGAAATGTAAATGATTAACAATCAAAATCATTATTCCT-----

137Lvolskensis TGGAACTTGTAAATGATTAACAATCAAAATAATCATTTCGGACTGAAAACCTTCCAAAAGGATATAAAACGTTTC

139Lholstii TGGAACTTGTAAATGATTAACAATCAAAATAATCATTTCGGACTGAAAACCTTCCAAAAGGATATAAAACGTTTC

143Lorientale CGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTAAAAAGGATATAAAATCTTTC

146Lnervosum TGGAACTTGTTCGGATTAACAATCCGAACCATTTAT-----

148Lhypericifoli TGGAAATGTAAATGATTAATAATCCGAAT-----GAAT-CACATTAT-----

149Lflavum CGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTAAAAAGGATATAAAATGTTTC

151Lcatharticum TGGAAATGTAAATGATTAACAATCAAAATAATTATTTGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

156Ltenuifolium TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAAACCTTACAAAAGGATATAAAACGTTTC

157Ltauricum CGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAAACCTTAAAAAGGATATAAAATGTTTC

163Llelegans CGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTAAAAAGGATATAAAATGTTTC

164Larboreum CGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTAAAAAGGATATAAAATGTTTC

165Lrzedowskii TGGAAATGTAAAGGATTAACAATCCGAACCATTTAT-----

171Lrlinoides -----

183Lpuberulum TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGACATT-----

184Lvernale TGGAAATGTAAATGATTAACAATCAAAATCATTAT-----TTC

186Llundellii TGGAAATGTAAATGATTAACAATCAAAATCATTAT-----TTC

187Lcremnophyllu TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

193Lbrevifolium TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

194Lprostratum GGGAAAGGGAAGGATAAACAAGCAAAATCATTATTCGGACGAAA-CTTACAAAAGGATATAAAACGTTTC

195Lchamissonis -----AGGATATAAAACATTTTC

196Lerigeroides -----

200Lcarneum TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAAACCTTACAAAAGGATATAAAACGTTTC

201Lpringlei TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAAACCTTACAAAAGGATATAAAACGTTTC

206Cselaginoides TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAAACCTTACAAAAGGATATAAAACGTTTC

210bLmacraeinacr TGGAAATTTTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

215Lchamissonis -----AGGATATAAAACATTTTC

219Lscoparium TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

220Cselaginoides TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

221Lburkartii TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

223Lkingii TGGAAATTTTAATTTTAATGATTAACAATCAAAATCATTATTCGGACTGAAAACCTTAGGATATAAAACGTTTC

224Lmonogynum TGGAACTTGTAAAGGATTAACAATCCGAATCATTAT-----

228Lneomexicanum TGGAAATGTAAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAAACGTTTC

231Lcompactum TGGAAATGTAAATGATTAACAATCAAAA-----CAAA-----

232Laustrale	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCT-----
233Laristatum	TGGAAATTGGAATGATTAACAATCAAAATCATTATTCT-----
234Lrigidumberla	TGGAAATTGTAATGATTAACAATCAAAA-----CAAA-----
236Lrigidumberla	TGGAAATTGTAATGATTAACAATCAAAA-----CAAA-----
238Lpratense	TGGAAATTGTAAGGATTAACAATCAAAACCATTTAT-----
241Lrigidumberla	TGGAAATTGTAATGATTAACAATCAAAA-----CAAA-----
242Lrupestre	-----
246LrigidumKerr	TGGAAATTGTAATGATTAACAATCAAAA-----CAAA-----
253Lelongatum	TGGAAATTGTAATGATTAACAATCAAAA-----CAAA-----
256Lmedium	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
257Lmedium	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
258Lquadrifolium	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
259Lheterostylum	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
266Lcarteri	TGGAAATTGTAATGATTAACAATCAAAATCATTATT-----C
268Larenicola	TGGAAAGTGAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
269Lwestii	TGGAAATTGTAATGATTAACAATCAAAA-----AGGATATATAAACGTTTC
270Lorizabae	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
291Lrigidumfilif	TGGAAATTGTAATGATTAACAATCAAAATCATTATT-----C
293Lquatemalense	TGGAAATTGTAATGATTAACAATCCGAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
294Lcatharticum	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
302Lsubteres	TGGAAATTGTAATGATTAACAATCAAAATCATTATT-----C
307Llalatium	TGGAAATTGTAATGATTAACAATCAAAATCATTATT-----C
310Llasiocarpum	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATAAACGTTTC
244Lhudsonioides	-----
214Lbienne	TGGAACTTGAAGGATTAACAATCCGAATCATTAT-----
249LlinumBrackenr	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCT-----
250Lhudsonioides	TGGAAATTGTAATGATTAACAATCAAAA-----
Al1Lrigidumpuberu	TGGAAATTGTAATGATTAACAATCAAAATCATTATT-----C
F1Lbahamensecora	TGGAAAGTGAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
F2Lcruciata	-----
F5Lnelsonii	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
F6Ltenellum	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
F7Lscabrellum	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
L17Ltnuifolium	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
L29Lmaritimum	CGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTAAAAAGGATAGAAAACGTTTC
L4Lusitatissimum	TGGAACTTGAAGGATTAACAATCCGAATCATTAT-----
L41Lviscosum	CGGAAATTGTAATGATTAATATCCGAAAT-----GAAT-CACCATTA-----
L43Lcatharticum	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
L57Lflavum	CGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTAAAAAGGATATATAAACGTTTC
LPB2Loligophyllu	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
PM1983Sdigynum	TGGAAATTGTAATGATTAACAATCCAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA1Lafricanum	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA10Lesterhuysen	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA11LinumAgulhas	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA12Laethiopicum	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA13LinumspWindb	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA14Lvillosum	-----
SA17Lthesioides	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA19Lcomptonii	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA3Lacuticarpum	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA4Lgracile	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA5Lgracile	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA6Lgracile	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA7Linumdehoop	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA8Ladustum	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SA9Lesterhuysena	TGGAAATTGTAATGATTAACAATCAAAATCATTATTCGGACTGAAA-CTTACAAAAGGATATATAAACGTTTC
SD1Lrigidumcompa	TGGAAATTGTAATGATTAACAATCAAAATCATTATT-----C
Gyp1YesoHills	TGGAAATTGTAATGATTAACAATCAAAATCATTATT-----C
400Lrigidumrigid	TGGAAATTGTAATGATTAACAATCAAAATCATTATT-----C
162Ldecumbens	TGGAACTTGAAGGATTAACAATACGAATCATTAT-----
401Lintercursum	-----

	22
	666666666666666666666666666666666677
	666667777777778888888888999999900000000011111111222222222223333333
Taxon/Node	5678901234567890123456789012345678901234567890123456789012345678901234567890123456
-----	-----
001Lstriatum	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
005Limbricatum	----ATTGG-AAAAAACTTTGAAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
006Lbahamense	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
007Lbahamense	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
009LSchiedeanum	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
010Lewisii	-----CTTGAAAAGTTTCGATCTTTTTT-T-----AGA-CCAATC
011Lsuffruticosu	TGACATTGT-AAAAAACCTTTAAATATTTTCGCACCTTTTGT-TTAATTGAC-----ATAGACCCAATC
012Lstrictum	TGACATTGT-AAAAAACCTTTAAATGTTCGCACCTTTTGT-TTAATTGAC-----ATAGACCCAATC
013Lnarbonense	-----CTTGCAAAGTTTCGATCAATTTT--TAATTGAC-----ATAGACCCAATC
022Hcleveandiii	TGACATTGG-AAAAAACCTTTGAATGTTCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
025Hdisjunctum	TGACATTGG-AAAAAACCTTTGAATGTTCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
027Lberlandierib	-----
028Lberlandierif	TGACATTGG-AAAAAACCTTTTACATGTTTCGCACCTTTTTT--TAATTGAC-----ATAGACCCAATC
032Lhdsonioides	-----
034Lrupestre	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
035Lrigidumrigid	TGACATTGG-AAAAAACCTTTTACATGTTTCGCACCTTTTTT--TAATTGAC-----ATAGACCCAATC
036Lrupestre	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
037Lmexicanum	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
038Llongipes	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
040Lcruciata	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
041Laristata	TGACATTGG-AAAAAACCTTTACATGTTTCGCACCTTTTTT--TAATTGAC-----ATAGACCCAATC
042Lflagellare	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
044Reinwardtiaia	TGACATTGG-AAAAAACCTGCAAAATGTTT-CACCTTTTTT--TAATTGAC-----ATAGACCCAATC
047Hdymarioides	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
048Mmicranthum	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
049Cselaginoides	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
055Laethiopium	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
059Lcomptonii	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
061Lesterhuysena	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
066Ltenuae	TAACATTGG-AAAAAACCTTTGAATGTTTTCGCACCTTTTTT-TTAATTGAG-----GTAGACCCAATC
062Lheterostylum	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
067Lthunbergii	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-T-----AATTGACATAGACCACAATC
072Ltrigynum	TAACATTGG-AAAAAACCTTTGAATGTTT-GACCTTTTTT-TTAATTGAC-----ATAGACCCAATC
075Lstriatum	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
076Lmarginale	-----CTTGCAAAGTTTCGATCAATTTT--TAATTGAC-----ATAGCCCAATC
083Sdigynum	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
089Llittorale	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
091Lloligophyllum	TGACATTGG-ACAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
123Lkingii	GGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
124Lflorianum	TGACCTTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
130Lvarginianum	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
132Lsulcatum	-GACATTGG-AAAAAACCTTTGAATGCTTCGCACCTTTTTT--TAATTGACTTTAATTGACATAGACCACAATC
137Lvolskensi	TGACATTGT-AAAAAACCTTTAAATGCTTCGCACCTTTTGT-TTAATTGAC-----ATAGACCCAATC
139Lholstii	TGACATTGG-AAAAAACCTTTAAATGCTTCGCACCTTTTGT-TTAATTGAC-----ATAGACCCAATC
143Lorientale	TAACATTGG-AAAAAACCTTTAAATGCTTCGCACCTTTTTT-TTAATTGAC-----ATAGACCCAATC
146Lnervosum	-----CTTGAAAAGTTTCGATCTTTTTT-T--TAATTGAC-----ATAGACCCAATC
148Lhypericifoli	-----CTTGAAAAGTTTCGACCCCTTTTT--TAATTG- -TTAATTGACATAGACCACAATC
149Lflovum	TAACATTGG-AAAAAACCTTTAAATGCTTCGCACCTTTTTT-TTAATTGAC-----ATAGACCCAATC
151Lcatharticum	TGACATTGG-AAAAAACCTTTAAATGCTTCGCACCTTTTTT--TAATTGAC-----ATAGACCCAATC
156Ltenuifolium	TGACATTGT-AAAAAACCTTTAAATGTTTCGCACCTTTTGT-TTAATTGAC-----ATAGACCCAATC
157Ltauricum	TAACATTGG-AAAAAACCTTTAAATGCTTCGCACCTTTTTT-TTAATTGAC-----ATAGACCCAATC
163Lelegans	TAACATTGG-AAAAAACCTTTAAATGCTTCGCACCTTTTTT-TTAATTGAC-----ATAGACCCAATC
164Larboreum	TAACATTGG-AAAAAACCTTTAAATGCTTCGCACCTTTTTT-TTAATTGAC-----ATAGACCCAATC
165Lzredowskii	-----CTTGAAAAGTTTCGATCTTTTTT-T-----AGA-CCAATC
171Rlinoids	-----TCGACCTTTTT--TATTGAC-----ATAGACCCAATC
183Lpuberulum	-----GG-AAAAAACCTTTTACATGTTTCGCACCTTTTTT--TAATTGAC-----ATAGACCCAATC
184Lvernale	TGACATTGG-AAAAAACCTTTTACATGTTTCGCACCTTTTTT--TAATTGAC-----ATAGACCCAATC
186Llundelli	TGACATTGG-AAAAAACCTTTTACATGTTTCGCACCTTTTTT--TAATTGAC-----ATAGACCCAATC
187Lcremnophyllu	TGACATTGG-ACAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
193Lbrevifolium	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGAC-----ATAGACCCAATC
194Lprostratum	TGACATTGG-ACAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
195Lchamissonis	TGACATTGG-AAAAAACCTTTGAATGTTCGCACCTTTTTT-TTAATTGACTTTAATTGACATAGACCACAATC
196Leriagroides	

200Lcarneum TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
201Lpringlei TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
206Cselaginoides TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
210bLmacraeimacr TGACATTGG-ACAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
215Lchamissonis TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
219Lscoparium TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGAC-----ATAGACCCAATC  
220Cselaginoides TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACA-----  
221Lburkartii TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
223Lkingii GGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
224Lmonogynum -----CTTGACAAAGTTTCGATCAATTTT--TAATTGAC-----ATAGCCCAATC  
228Lneomexicanum GGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
231Lcompactum -GACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
232Laustrale -GACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
233Laristatum -GACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
234Lrigidumberla -GACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
236Lrigidumberla -GACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
238Lpratense -----CTTGACAAAGTTTCGATCTTTTT--TAGA--CCAATC  
241Lrigidumberla -GACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
242Lrupestre -----  
246LrigidumKerr -GACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
253Lelongatum -GACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
256Lmedium TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
257Lmedium TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
258Lquadrifolium TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
259Lheterostylum TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
266Lcarteri TGACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
268Larenicola TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
269Lwestii TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
270Lorizabae TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAG-----  
291Lrigidumfilif TGACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
293Lguatemalense TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
294Lcatharticum TGACATTGG-AAAAAATTTTAAATGCTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
302Lsubteres TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT--TAATTGACTTTAATTGACATAGACCCAATC  
307Llalum TGACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
310Llasiocarpum TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
244Lhudsonioides -----  
214Lbienne -----CTTGACAAAGTTTCGATCAATTTT--TAATTGAC-----ATAGCCCAATC  
249LinumBrackenr -GACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
250Lhudsonioides -GACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT--TAATTGACTTTAATTGACATAGACCCAATC  
AlLrigidumpuberu TGACATTGG-AAAAAATTTTACATGTTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
FlLbahamensecora TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
F2Lcruciata -----  
F5Lnelsonii TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAAT-  
F6Ltenellum TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
F7Lscabrellum TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
L17Ltenuifolium TGACATTGT-AAAAAATTTTAAATGTTTCGACCTTTTGT-TTAATTGAC-----ATAGACCCAATC  
L29Lmaritimum TAACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGAC-----ATAGACCCAATC  
L4Lusitatissimum -----CTTGACAAAGTTTCGATCAATTTT--TAATTGAC-----ATAGCCCAATC  
L41Lviscosum -----CTTGACAAAGTTTCGACCCCTTTT--TAATTG--TTTAATTGACATAGATCCAATC  
L43Lcatharticum TGACATTGG-AAAAAATTTTAAATGCTTCGACCTTTTTT--TAATTGAC-----ATAGACCCAATC  
L57Lflavum TAACATTGG-AAAAAATTTTAAATGCTTCGACCTTTTTT-TTAATTGAC-----ATAGACCCAATC  
LPB2Loligophyllu TGACATTGG-ACAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
PM1983Sdigynum TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTTTTAAATTGACTTTAATTGACATAGACCCAATC  
SA1Lafricanum TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA10Lesterhuysen TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA11LinumAgulhas TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA12Laethiopicum TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA13LinumspWindb TGACATTGG-AAAAAATTTTGAAATGTTTGACCTTTTTT--TAATTGACTTTAATTGACATAGACCCAATC  
SA14Lvillosum -----  
SA17Lthesioides TGACTTTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA19Lcomptonii TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA3Lacuticarpum TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA4Lgracile TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA5Lgracile TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA6Lgracile TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA7Linumdehoop TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA8Ladustum TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC  
SA9Lesterhuysena TGACATTGG-AAAAAATTTTGAAATGTTTCGACCTTTTTT-TTAATTGACTTTAATTGACATAGACCCAATC



SD1Lrigidumcompa TGACATTGG-AAAAAACTTTTACATGTTTCGACCTTTTT--TAATTGAC-----ATAGACCCAATC  
 Gyp1YesoHills TGACATTGG-AAAAAACTTTTACATGTTTCGACCTTTTT--TAATTGAC-----ATAGACCCAATC  
 400Lrigidumrigid TGACATTGG-AAAAAACTTTTACATGTTTCGACCTTTTT--TAATTGAC-----ATAGACCCAATC  
 162Ldecumbens -----CTTGACAAAGTTTCGATCAATTTT--TAATTGAC-----ATAGACCCAATC  
 401Lintercursum -----

222222222222  
 777777777777  
 333444444444  
 Taxon/Node 789012345678  
 -----

001Lstriatum CATTTAGTAAAG  
 005Limbricatum CATTTAGTAAAG  
 006Lbahamense CA-----  
 007Lbahamense CATTTAGTAAAG  
 009Lschiedeanum CATTTAGTAAAG  
 010Llewisii CGTTTACTAAAG  
 011Lsuffruticosu CATTTAGTAAAG  
 012Lstrictum CATTTAGTAAAG  
 013Lnarbonense CATTTAGTAAAG  
 022Hclevelandii CATTTAGTAAAG  
 025Hdisjunctum CATTTAGTAAAG  
 027Lberlandierib -----  
 028Lberlandierif CATTTAGTAAAG  
 032Lhudsonioides -----  
 034Lrupestre CATCTAGTAAAG  
 035Lrigidumrigid CATTTAGTAAAG  
 036Lrupestre CATYTAGTAAAG  
 037Lmexicanum CATTTAGTAAAG  
 038Llongipes CATTTAGTAAAG  
 040Lcruciata CATTTAGTAAAG  
 041Laristatum CATTTAGTAAAG  
 042Lflagellare CATTTAGTAAAG  
 044Reinwardtiaia CATTTAGTAAAG  
 047Hdrymarioides CATTTAGTAAAG  
 048Hmicranthum CATTTAGTAAAG  
 049Cselaginoides CATTTAGTAAAG  
 055Laethiopicum CATTTAGTAAAG  
 059Lcomptonii CATTTAGTAAAG  
 061Lesterhuysena CATTTAGTAAAG  
 066Ltenue CATTTAGTAAAG  
 062Lheterostylum C-TTTAGTAAAG  
 067Lthunbergii CATTTAGTAAAG  
 072Ltrigynum CATTTAGTAAAG  
 075Lstriatum CATTTAGTAAAG  
 076Lmarginale CATTTAGTAAAG  
 083Sdigynum CATTTAGTAAAG  
 089Llittorale CATTTAGTAAAG  
 091Loligophyllum CATTTAGTAAAG  
 123Lkingii CATTTAGTAAAG  
 124Lfloridanum CATTTAGTAAAG  
 130Lvirginianum CATTTAGTAAAG  
 132Lsulcatum CATTTAGTAAAG  
 137Lvolkensii CATTTAGTAAAG  
 139Lholstii CATTTAGTAAAG  
 143Lorientale CATTTAGTAAAG  
 146Lnervosum CGTTTCGTAAAG  
 148Lhypericifoli CGTTTAGTAAAG  
 149Lflavum CATTTAGTAAAG  
 151Lcatharticum CATTTAGTAAAG  
 156Ltenuifolium CATTTAGTAAAG  
 157Ltauricum CATTTAGTAAAG  
 163Lelegans CATTTAGTAAAG  
 164Larboreum CATTTAGTAAAG  
 165Lrzedowskii CGTTTACTAAAG  
 171Rlinoides CATTTAGTAAAG  
 183Lpuberulum CATTTAGTAAAG

184Lvernale	CATTTAGTAAAG
186Llundellii	CATTTAGTAAAG
187Lcremnophyllu	CATTTAGTAAAG
193Lbrevifolium	CATTTAGTAAAG
194Lprostratum	C-TTTAGTAAA-
195Lchamissonis	CATTTAGTAAAG
196Lerigeroides	-----
200Lcarneum	CATTTA-----
201Lpringlei	CATTTAGTAAAG
206Cselaginoides	CATTTAGTAAAG
210bLmacraeimacr	CATTTAGTAAAG
215Lchamissonis	CATTTAGTAAAG
219Lscoparium	CATTTAGTAAAG
220Cselaginoides	-----
221Lburkartii	CATTTAGTAAAG
223Lkingii	CATTTAGTAAAG
224Lmonogynum	CATTTAGTAAAG
228Lneomexicanum	CATTTAG-----
231Lcompactum	CATTTAGTAAAG
232Laustrale	CATTTAGTAAAG
233Laristatum	CATTTAGTAAAG
234Lrigidumberla	CATTTAGTAAAG
236Lrigidumberla	CATTTAGTAAAG
238Lpratense	CGTTTACTAAAG
241Lrigidumberla	CATTTAGTAAAG
242Lrupestre	-----
246LrigidumKerr	CATTTAGTAAAG
253Lelongatum	CATTTAGTAAAG
256Lmedium	CATTTAGTAAAG
257Lmedium	CATTTAGTAAAG
258Lquadrifolium	CATTTAGTAAAG
259Lheterostylum	CATTTAGTAAAG
266Lcarteri	CATTTAGTAAAG
268Larenicola	C-----
269Lwestii	CATTTAGTAAAG
270Lorizabae	-----
291Lrigidumfilif	CATTTAGTAAAG
293Lquatemalense	CATTTAGTAAAG
294Lcatharticum	CATTTAGTAAAG
302Lsubteres	CATTTAGTAAAG
307Lalatum	CATTTAGTAAAG
310Llasiocarpum	CATTTAGTAAAG
244Lhudsonioides	-----
214Lbienne	CATTTAGTAAAG
249LinumBrackenr	CATTTAGTAAAG
250Lhudsonioides	CATTTAGTAAAG
A1Lrigidumpuberu	CATTTAGTAAAG
F1Lbahamensecora	CATTTAGTA---
F2Lcruciata	-----
F5Lnelsonii	-----
F6Ltenellum	CATTTAGTAAAG
F7Lscabrellum	CATTTAGTAAAG
L17Ltenuifolium	CATTTAGTAAAG
L29Lmaritimum	CATTTAGTAAAG
L4Lusitatissimum	CATTTAGTAAAG
L41Lviscosum	CGTTTAGTAAAG
L43Lcatharticum	CATTTAGTAAAG
L57Lflavum	CATTTAGTAAAG
LPB2Loligophyllu	CATTTAGTAAA-
PM1983Sdigynum	CATTTAGTAAAG
SA1Lafricanum	CATTTAGTAAAG
SA10Lesterhuysen	CATTTAGTAAAG
SA11LinumAgulhas	CATTTAGTAAAG
SA12Laethiopicum	CATTTAGTAAAG
SA13LinumspWindb	CATTTAGTAAAG
SA14Lvillosum	-----
SA17Lthesioides	CATTTAGTAAAG
SA19Lcomptonii	CAT-----

SA3Lacuticarpum	CATTTAGTAAAG
SA4Lgracile	CATTTAGTAAAG
SA5Lgracile	CATTTAGTAAAG
SA6Lgracile	CATTTAGTAAAG
SA7Linumdehoop	CATTTAGTAAAG
SA8Ladustum	CATTTAGTAAAG
SA9Lesterhuysena	CATTTAGTAAAG
SD1Lrigidumcompa	CATTTAGTAAAG
Gyp1YesoHills	CATTTAGTAAAG
400Lrigidumrigid	CATTTAGTAAAG
162Ldecumbens	CATTTAGTAAAG
401Lintercursum	-----

## Appendix D: Selected ITS Matrix, Chapter 3

[illegible]

049Cselaginoides -----CATTGTGCGAAACCTGC-AAAGCA  
206Cselaginoides -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
210Lmacraei -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
210Lmacraei -----CAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
215Lchamissonis -----CGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
219Lscoparium -----AACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
220Cselaginoides -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
221Lburkartii -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
223Lkingii -----AACCTGC-AAAGCA  
228Lneomexicanum -----TAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
231Lcompactum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
232Laustale -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
233Laristatum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
234Lberlandieri -----CAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
236Lrigidum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
241Lberlandieri -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
242Lrupestre -----TAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
244Lhudsonioides -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
246LrigidumKerr -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
249Limbristatum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
250Lhudsonioides -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
251Lrigidum -----TGTCGAAACCTGC-AAAGCA  
253Lelongatum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
256Lmedium -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
257Lmedium -----AGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
258Lquadrifolium ---GGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
266Lcarteri -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
268Larenicola -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
269Lwestii -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
270Lorizabae -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
291Lrigidumfilif -----GTGCGAAACCTGC-AAAGCA  
293Lguatemalense -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
294Lcatharticum -----TCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
307Llalatium GGAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
310Llasiocarpum -----GTGCGAAACCTGC-AAAGCA  
401Lintercursum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
AlLrigidum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
FlLbahamensecora ---GGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
F2Lcruciata -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
F5Lnelsonii -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
F6Ltenellum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
F7Lscabrellum -----GATCATTGTGCGAAACCTGC-AAAGCA  
F8Ltenellum -----GTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
L17Ltenuifolium -----CATTGTGCGAAACCTGC-AAAGCA  
L29Lmaritimum -----CATTGTGCGAAACCTGC-AAAGCA  
L43Lcatharticum -----CATTGTGCGAAACCTGC-AAAGCA  
L57Lflavum -----CATTGTGCGAAACCTGC-AAAGCA  
LinumSpYeso -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
PM1983Sdigynum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA10Lesterhuysen -----AGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA11LinumAgulhas -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA12Laethiopicum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA13LinumspWindb -----TAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA14Lvillosum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA17Lthesiodes -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA19Lcompton -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA1Lafricanum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA3Lacuticarpum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA4Lgracile -----TGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA5Lgracile -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA6Lgracile -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA7LinumDeHoop -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA8adustum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA8Ladustum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SA9Lesterhuysena -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA  
SDLinum -GAAGGAGAAGTCGTAACAAGGTTTCCGTAGGTGAACCTGCGGAAGGATCATTGTGCGAAACCTGC-AAAGCA

[illegible]

221Lburkartii	GAATGACC-AGCGAACTCGTTGTGTTA---AACGTCGAACGGGGTCGGAGCGAGATGCGGCCCTTT-GGCCCTCC
223Lkingii	GAATGACC-AGAGAACTCGTTGTGTTA---AATGTCC-ACGGGGTCGGAGCGAGTTGTGGCCATT-GGCCCTCC
228Lneomexicanum	GAATGACC-AGTGAACCTGTTGTGTTA---AATGTCG-ACGGGGTCGGAGCGAGTTGTGGCCATT-GGCCCTCC
231Lcompactum	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
232Laustrale	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
233Laristatum	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
234Lberlandieri	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
236Lrigidum	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
241Lberlandieri	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
242Lruepreste	GAATGACC-AGCGAACTCGTTGTGTTA---AATGTCG-ACGGGGTCGGAGCGAGTTGTGGCCCTTC-GGCCCTCC
244Lhudsonioides	GAATGACC-AGCGAACTCGTTGTGTTA---AAAGTCG-ACGGGGTCGGAGCGAGTTGTGGCCCTTC-GGCCCTCC
246LrigidumKerr	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
249Limbriatum	GAATGACC-AGCGAACTCGTTGTGTTA---AATGTCG-ACGGGGTCGGAGCGAGTTGTGGCCCTTC-GGTCTCC
250Lhudsonioides	GAATGACC-AGCGAACTCGTTGTGTTA---AAAGTCG-ACGGGGTCGGAGCGAGTTGTGGCCCTTC-GGCCCTCC
251Lrigidum	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
253Lelongatum	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
256Lmedium	GAATGACC-AGCGAACTCGTTGTGTTA---AATGTCG-ATGGGGTCAGAGCGAGTTGTGGCCCTTC-GGCCCTCC
257Lmedium	GAATGACC-AGCGAACTCGTTGTGTTA---AATGTCG-ATGGGGTCAGAGCGAGTTGTGGCCCTTC-GGCCCTCC
258Lquadrifolium	GAATGACC-AGCGAACTTGTAGTGTGTTA---AATGTCG-ATGGGTGTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC
266Laricaria	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
268Larenicola	GAATGACC-AGCGAACTCGTTGTGTTA---AATGTCG-ACGGGGTCGGAGCGAGTTGTGGCCCTTC-GGCCCTCC
269Lwestii	GAATGACC-TGCGAACTCGTTGTGTTA---AATGTCG-ACGGGGTCAGAGCGAGTTGTGGCCCTTC-GGCCCTCC
270Lorizabae	GAATGACC-AGTGAACCTCGTTGTGTTA---AACGTCG-ACGGGGTCGGAGCGAGATGCGGCGTCT-AGCCCTCC
291Lrigidumfilif	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
293Lguatemalense	GAATGACC-AGTGAACCTCGTTGTGTTA---AACGTCG-ACGGGGTCGGAGCGAGATGCGGCGTCT-AGCCCTCC
294Lcatharticum	GAACGACC-AGCGAACTCGTTGTGTTA---AACTTTG-CGCGGGTCGGAGCTAGATGCGGCCCTTC-GGCCCTTC
307Lalatum	GAATGACC-AGCGAACTCGTTGTGTTA---A-CTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
310Llasiocarpum	-----ACTCGTTGTGTTA---AATGTTG-ACGGGGTCAGAGCGAGTATGCGGCCCTTC-GGCCCTCC
401Lintercursum	GAATGACC-AGCGAACTCGTTGTGTTA---AATGTCG-ATGGGGTCAGAGCGAGTTGTGGCCCTTC-GGCCCTCC
AlLrigidum	GAATGACC-AGCGAACTCGTTGTGTTA---AACTTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
F1Lbahamensecora	GAATGACC-AGCGAACTTGTGTGTTA---AATGTCG-ACGGGGTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC
F2Lcruciata	GAATGACC-AGCGAACTTGTGTGTTA---AATGTCG-ATAGGGTCCGAGCGAGTGTGGCCCTTC-GGCCCTCC
F5Lnelsonii	GAATGACC-AGCGAACTTGTGTGTTA---AATGTTG-ATAGGGTCCGAGCGAGTGTGGCCCTTC-GGCCCTCC
F6Ltenellum	AAATGACC-AGTGAACCTCGTTGTGTTA---AATGTTG-ACGAGGTTCAGAGCGAGTGTGGCCCTTC-GGCCCTCC
F7Lscabrellum	GAATGACC-AGCGAACTCGTTGTGTTA---AATGTCG-ACGGGGTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC
F8Ltenellum	GAATGACC-AGCGAACTTGTGTGTTA---AATATCA-ACGGGGTCAAAGCGAGTGTGGCCCTTC-GGCCCTCC
L17Ltenuifolium	GAATGACC-AGCGAACTTGTGTGTTA---AACTTCG-CGCGGGCTAG-----CTCT
L29Lmaritimum	GAATGACC-AGCGAACTCGTTGTGTTA---AACATGG-ACAAGGTCAGAGCGAGATGTGGCCCTTG-GCCATCT
L43Lcatharticum	GAACGACC-AGCGAACTCGTTGTGTTA---AACTTTG-CGCGGGTCGGAGCTAGATGCGGCCCTTC-GGCCCTTC
L57Lflavum	GAATGACC-AGCGAACTCGTTGTGTTA---ATCTCTG-ACGGGGTCGGACCGAGTGTGGCCCTTC-GTCTCT
LinumSpYeso	GAATGACC-AGCGAACTCGTTGTGTTA---AATCTCG-ACGGGGTCGGACCGAGCGGTGGCCCTTC-GGGCTCC
PM1983Sdigynum	GAATGACC-AGCGAACTCGTTGTGTTA---AATGTCG-ACGAGTTCGGAGCGAGGAGCGGCTCTGGCCCTTC
SA10Lesterhuysen	GAATGACC-AGCGAACTCGTAGTGTGTTA---AATGTCG-ATGGGGTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC
SA11LinumAguilhas	GAATGACC-AGCGAACTCGTTGTGTTA---AATGTTG-ACGGGGTCATAGCGAGTGTGGCCCTTC-GGCCCTCC
SA12Laethiopicum	GAATGACC-AGCGAACTTGTAGTGTGTTA---AATGTCG-ACGGGGTCATAGCGAGTGTGGCCCTTC-GGCCCTCC
SA13LinumspWindb	GAATGACC-AGCGAACTTGTAGTGTGTTA---AATGTCG-ACGGGGTCATAGCGAGTGTGGCCCTTC-GGCCCTCC
SA14Lvillosum	GAATGACC-AGCGAACTCGTAGTGTGTTA---AATGTCG-ACGGGGTCAGAGCGAGTGTGGCCCTTC-GGCCCTCC
SA17Lthesiodes	GAATGACC-AGCGAACTTGTAGTGTGTTA---AATGTCG-ACGGGGTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC
SA19Lcompton	GAATGACC-AGCGAACTCGTAGTGTGTTA---AATGTCG-ACGGGGTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC
SA1Lafricanum	GAATGACC-AGCGAACTCGTAGTGTGTTA---AATGTCG-ACGGGGTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC
SA3Lacuticarpum	GAATGACC-AGCGAACTCGTAGTGTGTTA---AATGTCG-CGCGGGTTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC
SA4Lgracile	GAATGACC-AGCGAACTTGTAGTGTGTTA---AATGTCG-ACGGGGTTCGGAGCAAGCTGTGGCCCTTC-GGCCCTCC
SA5Lgracile	GAATGACC-AGCGAACTTGTAGTGTGTTA---AATGTCG-ACGGGGTCATAGCGAGTGTGGCCCTTC-GGCCCTCC
SA6Lgracile	GAATGACC-AGCGAACTTGTAGTGTGTTA---AATGTCG-ATGGGGTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC
SA7LinumDeHoop	GAATGACC-AGCGAACTTGTAGTGTGTTA---AATGTCG-ACGGGGTCATAGCGAGTGTGGCCCTTC-GGCCCTCC
SA8adustum	GAATGACC-AGCGAACTTGTAGTGTGTTA---AATGTCG-ACGGGGTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC
SA8Ladustum	GAATGACC-AGCGAACTTGTAGTGTGTTA---GATGTCG-ACGGGGTCATAGCGAGTGTGGCCCTTC-GGCCCTCC
SA10Lesterhuysen	GAATGACC-AGCGAACTCGTAGTGTGTTA---AATGTCG-ATGGGGTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC
SDLinum	GAATGACC-AGCGAACTCGTTGTGTTA---AATGTCG-ACGGGGTCGGAGCGAGTGTGGCCCTTC-GGCCCTCC

007Lbahamense TCGAGATGGCCACGTCGGGC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 009Lschiedeanum ATGCAATGGCCTTATCAGTG-GGGG---GATCGGCCCTAGTGGTT--GTTCC-TTCCCGCACAACT-AACA  
 011Lsuffruticosu ATGCGATGGCCTCGTCGGGT---GG-GGGCTCGGCCCTCGTGGTC--GTGCC--TTCCCGCACAACT-AACA  
 012Lstrictumssps ACGCGAAGGCCTCGTTGGGT--GGT-GGGCTCGGCCCTTGTGGCC--TTGCC-TTCCACACAACT-AACA  
 022Hclevelandii GCGCAATGGCCTCGTCGAGCGGGG---GCTTGGCCCTTGTGGTC--GTGCC-TTCTGCACAACT-AACA  
 025Hdisjunctum GCGCAATGGCCTCGTCGAGCGGGG---GCTTGGCCCTTGTGGTC--GTGCC-TTCTGCACAACT-AACA  
 027Lberlandierib TCGCGATGGCCACGTCGTGC-GGTG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 028Lberlandierif TCGCGATGGCCACGTCGTGC-GGGG---GCTTGGCCCTTGTGGTC--GTTCC-TTCCCGCRCAAACT-AACA  
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 034Lrupestre TCGCGATGGCCACGTCGGGC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 035Lrigidum TCGCGATGGCCACGTCGTGC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 036Lrupestre TCGCGATGGCCACGTCGGGC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 037Lmexicanum ACGCGATGGCCTCGTCGAGC-GGGG---GCTTGGCCCTTGTGGTC--GTGCC-TTCCCGCACAACT-AACA  
 038Llongipes ACGTGATGACCTCGTCGAGC-GGGG---GCTCAGCC-TTGTGGTT--GTGCC-TTCCCGCACAACT-AACA  
 040Lcruciata ACGTGATGACCTCGTCGAGC-GGGG---GCTCAGCC-TTGTGGTT--GTGCC-TTCCCGCACAACT-AACA  
 041Laristatum TCGTGATGGCCGTCGTGC-GGGG---GCTTGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 042Lflagellare TCGCGATGGCCACGTCGGGC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 047Hdrymarioides ATGTGATGGCCTCGTCGAGC-GGGA---GCTTGGCCCTTGTGGTC--GTGCC-TTCTGCACAACT-AACA  
 048Hmicranthum GCGCAATGGCCTCGTCGAGC-GGGA---GCTTGGCCCTTGTGGTC--GTGCC-TTCTGCACAACT-AACA  
 055Laethiopicum ACGCGATGGCCTCATCGGGC-GGGG---GCTTGACTCTCGTGGTC--GTGCC-TTCCCGCACAACT-AACA  
 059Lcomptonii ACGCGATGGCCTCGTCGAGC-GGGG---GCTTGACTCTCGTGGTC--GTGCC-TTCCCGCACAACT-AACA  
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 062Lheterostylum ACGCGATGGCCTCGTCGGGC-GGGG---GCTTGACTCTCGTGGTC--ATGCC-TTCCCGCACAACT-AACA  
 066Ltenue TAGTGATGACCTCGTCTAGA-GGGG---TGCTGGTCTAGCGGCC--ATTCT-TTCCCGCACAACT-AACA  
 067Lthunbergii ACGTGATGGCCTCGTCGGGC-GGGG---GCTTGATCTCGTGGTC--GTGCC-TTCCCGCACAACT-AACA  
 072Ltrigynum TAGTGATGACCTCGTCTAGA-GGGG---TGCTGGTCTAGTGGCC--GTTCT-TCCCTCAAAACT-AACA  
 075Lstriatum ACGTTATGGCCTCGTCGGGC-GGAG---GCTTGGCCCTCGTGGCC--TTGCC-TACCCGCACAACT-AACA  
 083Sdigynum AAGCGACGACCTCGTCGAGCGGTG---GCTCGGCCCTTGTGGTC--GTGCC-TTCCCGCACAACT-AACA  
 089Llittorale ACGCGATGGCCTTGTGGGC-GGGG---GCTCAGCCGTCGTGGTT--GTTCC-CTCCCGCACAACT-AACA  
 091Lloliophyllum ACGCGATGGCCTCGTCGGGC-GGGG---GCTCAGCCGTCGTGGTT--GTGCC-CTYCCGCACAACT-AACA  
 123Lkingii TCGCGATGGCCTTGTGGGC-GGGG---GCTCAGCCCTCGTGGTC--GTGCM-TTCCCGCACAACT-AACA  
 124Lfloridanum ACGTTATGGCCTCGTCGGGC-GGAG---GCTTGGCCCTCGTGGCC--TTGCC-TACCCGCACAACT-AACA  
 130Lvirginianum ACGTTATGGCCTCGTCGGGC-GGAG---GCTTGGCCCTCGTGGTC--TTTGCC-TACCCGCACAACT-AACA  
 132Lsulcatum CCGCAATGGCCACGTCGGCGGGG---GCATGGCCCTTGTGGTC--ACTCC-TTCCCGCACAACT-AACA  
 137Lvolkensii ACGCGATGGCCTCGTCGGGT--GGT-GGGCTTGGTCTTGTGGCC--GTGCC-TTCCCGCACAACT-AACA  
 139Lholstii ACGCGATGGCCTCGTCGGGT--GGT-GGGCTTGGTCTTGTGGCC--GTGCC-TTCCCGCACAACT-AACA  
 143Lorientale GTGTGATGGCCTCGTCTTGC-GGGATTGTCGCGTCTAGTGGCC--TTGCC-TTCCCGCACAACT-AACA  
 149Lflavum GTGTGATGGCCTCGTCTTGC-GGGATTGTCGCGTCTAGTGGCC--TTGCC-TTCCCGAACAACT-AACA  
 151Lcatharticum ATGCGATGGCCTCGTCGGGC---GT-GGGCTCGGTCTAGTGGCT--GTTCC-TTACCGCACAACT-AACA  
 156Ltenuifolium ATGCGATGGCCTCGTCGGGT-GGGG---GCTCGGCCCTCGTGGTC--GTGTC--TTCCCGCACAACT-AACA  
 157Ltauricum GTGTGATGGCCTCGTCTTGC-GGGATTGTCGCGTCTAGTGGCC--TTGCC-TTCCCGCACAACT-AACA  
 163Lelegans GTGTGATGGCCTCGTCTTGC-GGGATTGTCGCGTCTAGTGGCC--TTGCC-TTCCCGCACAACT-AACA  
 164Larbareum GTGTGATGGCCTCGTCTTGC-GGGATTGTCGCGTCTAGTGGCC--TTGCC-TTCCCGCACAACT-AACA  
 171Rlinoides TCGTGTCGAACCTCGTCGAGC-GGGT-GGGGTTGGTCTCGTGGCC--TTGCC-TTCCCGCACAACT-AACA  
 183Lpuberulum TCGTGATGGCCCGTCGTGC-GGGG---GCTTGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 184Lvernale TCGTGACGGCCACGTCGTGC-GGGG---GCTTGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 186Llundellii TCGTGATGGCCACGTCGTGC-GGTG---GCTTGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 187Lcremophyllu ACGCGATGGCCTCGTCGGGC-GGGG---GCTCAGCCGTCGTGGTT--GTGCC-CTCCCGCACAACT-AACA  
 193Lbrevifolium ACGCGATGGCCTCGTCGGGC-GGGG---GCTCAGCCCTCGTGGTT--GTGCC-TTCTGCACAACT-AACA  
 194Lprostratum ACGCGATGGCCTCGTCGGGC-GGGG---GCTCAGCCGTCGTGGTT--GTGCC-CTCCCGCACAACT-AACA  
 195Lchamissonis ACGCAATGGCCTCATCGGGC-GGGG---GCTCAGCCGTCGGGTT--GTGCC-CTCCACAAAACT-AACA  
 196Lerigeroides ACGCGATGGCCTCGTCGGGC-GGGG---GCTCAGCCGTCGTGGTT--GTGCC-CTCCCGCACAACT-AACA  
 200Lcarneum ACGCGATGGCCTCGTCGGGC-GGGG---GCTCAGCCGTCGTGGTT--GTGCC-CTCCCGCACAACT-AACA  
 201Lpringlei TCGCAATGGCCACGTCGGGC-GGGG---GCTTGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 049Cselaginoides ACGCGATGGCCTCGTCGGCGGGG---GCTCAGCCACGTGGTT--GTGCC-TTCCCGCACAACT-AACA  
 206Cselaginoides ACGCGATGGCCTCGTCGGCGGGG---GCTCAGCCACGTGGTT--GTGCC-TTCCCGCACAACT-AACA  
 210Lmacraei ACGCGATGGCCTCGTCGGGC-GGGG---GCTCAGCCGTCGTGGTT--GTGCC-CTCCCGCACAACT-AACA  
 210Lmacraei ACGCGATGGCCTCGTCGGGC-GGGG---GCTCAGCCGTCGTGGTT--GTGCC-CTCCCGCACAACT-AACA  
 215Lchamissonis ACGCAATGGCCTCATCGGGC-GGGG---GCTCAGCCGTCGTGGTT--GTGCC-CTCCACAAAACT-AACA  
 219Lscoparium ACGCGATGGCCTCGTGGCGGGG---ACTCAGCCCTCGTGGTT--GTGCC-TTCCCGCACAACT-AACA  
 220Cselaginoides ACGCGATGGCCTCGTCGGCGGGG---GCTCAGCCACGTGGTT--GTGCC-TTCCCGCACAACT-AACA  
 221Lburkartii ACGTGATGGCCTCGTAGGGC-GAGG---GCTCAGCCATGTGGTT--GTGCC-TTCCCGCACAACT-AACA  
 223Lkingii TCGCGATGRCCTTGTGGGC-GGGG---GCTCAGCCCTCGTGGTA--GTGCC-TTCCCGCACAACT-AACA  
 228Lneomexicanum CCGTGATGGCCTCGCGGGC-GGGG---GCTTACCCCTCGTGGTC--TTGCT-TTCCCGCACAACT-AACA  
 231Lcompactum TCGCGATGGCCACGTCGTGC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 232Laustrale TCGTGATGGCCACGTCGTGC-GGGG---GCTTGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 233Laristatum TCGTGATGGCCACGTCGTGC-GGGG---GCTTGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA  
 234Lberlandieri TCGCGATGGCCACGTCGTGC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAACT-AACA



236Lrigidum	TCGCGATGGCCACGTCGTGTC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
241Lberlandieri	TCGCGATGGCCACGTCGTGTC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
242Lrupestr	TCGCGATGGCCACGTCGGGC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
244Lhudsonioides	TCGCGACGCGCGCTCGTGTGTC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
246LrigidumKerr	TCGCGATGGCCACGTCGTGTC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
249Limbricatum	TCGTGATGGCCACGTCGTGTC-GGTG---GCTCGGCCCTTGTGGCC--GTTCC-TTCCCGCACAAACT-AACA
250Lhudsonioides	TCCGACGCGCGCTCGTGTGTC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
251Lrigidum	TCGCGATGGCCACGTCGTGTC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
253Leelongatum	TCGTGATGGCCACGTCGTGTC-GGGG---GCTTGGCCCTTGTGGTC--GTTCC-TCCCGCACAAACT-AACA
256Lmedium	ACGTTATGGCCTCGTCGGGC-GGAG---GCTTGGCCCTCGTGGTCTTTTGGC-TACCCGCACAAACT-AACA
257Lmedium	ACGTTATGGCCTCGTYGGGC-GGAG---GCTTGGCCCTCGTGGTCTTTTGGC-TACYCGCACAAACT-AACA
258Lquadrifolium	ACGCGATGGCCTCGTCGGGC-GGGG---GCTTGACTCTCGTGGTC--GTGCC-TTCCCGCACAAACT-AACA
266Lcarteri	TCGTGATGGCCACGTCGTGTC-GGGG---GCTTGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
268Larenicola	TCGCGATGGCCACGTCGGGC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
269Lwestii	ACGTATGCGCCTCGTCGGGC-GGAG---GCTTGGCCCTCGTGGTC--GTGCC-TACCCGCACAAACT-AACC
270Lorizabae	ACGCGATGGCCTCGTCGAGC-GGGA---GCTTGGCCCTTGTGGTC--GTGCC-TTCTGCACAAACC-AACA
291Lrigidumfilif	TCGCGATGGCCACGTCGTGTC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCACACAAACT-AACA
293Lguatemalense	ACGCGATGGCCTCGTCGAGC-GGGG---GCTTGGCCCTTGTGGTC--GTGCC-TTCTGCACAAACC-AACA
294Lcoarthicum	ATGCGATGGCCTCGTCGGGC---GT-GGGCTCGGTCTAGTGGCT--GTTCC-TTACGCCACCAACA-AACA
307Lalatum	TCGTGATGGCCACGTCGTGTC-GGGG---GCTTGGCCCTTGTGGTC--TTTCC-TTCCCGCACAAACT-AACA
310Llasiocarpum	ACGCGATGGCTCTGTTGGGC-AGGA---GCTCGGCCCTCGTGGTT--GTTGC-TTCCCGCACAAACT-AACA
401Lintercursum	ACGTTATGGCCTCGTCGGGC-GGAG---GCTTGGCCCTCGTGGTCTTTTGGC-TACCCGCACAAACT-AACA
AlLrigidum	TCGTGACGGCCACGTCGTGTC-GGGG---GCTTGGCCCTTGTGGTC--GCTCC-TTCCCGCACAAACT-AACA
F1Lbahamensecora	TCGAGATGGCCACGTCGGGC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
F2Lcruciata	ACGCAACGGCCTTGTCAGTG-GGGG---GATTGGCCCTAGTGGTT--GTTCC-TTCCCGCACAAACT-AACA
F5Lnelsonii	ATGCAATGGCCTTATCAGTG-GGGG---GATCGGCCCTAGTGGTT--GTTCC-TTCCCGCACAAACT-AACA
F6Ltenellum	ACGCGATGGCTCTGTTGGGC-AGGA---GCTCGGCCCTCGTGGTT--GTTGC-TTCCCGCACAAACT-AACA
F7Lscabellum	TCGCGATGGCCACGTCGGGC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
F8Ltenellum	ATGTGATGACCTCTGTTGGGTGGGG---GATCGGCCCTCGTGGTC--GTTCT-TTCCCGCACAAACT-AACA
L17Ltenuifolium	ATGCGAAGGCCTCGTCGGGT-GGGG---GCTCGGCCCTCGTGGTC--GTGCC--TTCCCGCACAACTAACA
L29Lmaritimum	TAGCGATGGCCCTGTCCGATGGGAT---GTTCGTCTAGTGGCY--GTTCC-TACCTCACAACACT-AACA
L43Lcoarthicum	ATGCGATGGCCTCGTCGGTC---GT-GGGCTCGGTCTAGTGGCT--GTTCC-TTACGCCACCAACA-AACA
L57Lflavum	TCGTGATGGCCTCGTCTTGC-GGATTTGCTCGTCTAGTGGCC--TTGCC-TTCCCGCACAAACT-AACA
LinumSpYeso	TCGTGATGGACCCGTCGCGC-GGGG---GCTTGGCCCTTGTGGTC--GTTCC-TCCCGCACAAACT-AACA
PM1983Sdigynum	AACGACACGCTCGTCGAGCGGTGG---GCTCGGCCCTTGCCTGTC--GTGCC-TTCCCGCACAAACT-AACA
SA10Lesterhuysen	ACGCGATGGCCTCCTCGGGC-GGGG---GCTTACTCTCGTGGTC--GTGCC-TTCCCGCACAAACT-AACA
SA11LinumAguilhas	ATCGATGGCCTCATCGGC-GGGG---GCTTACTCTCGTGGTC--GTGCC-TTCCCGCATAAACT-AACA
SA12Laethiopicum	ACGCGATGGCCTCATCGGC-GGGG---GCTTACTCTCGTGGTC--GTGCC-TTCCCGCATAAACT-AACA
SA13LinumspWindb	ACGCGATGGCCTCATCGGC-GGGG---GCTTACTCTCGTGGTC--GTGCC-TTCCCGCATAAACT-AACA
SA14Lvillosum	ACGCGATGGCCTTGTGCGGC-GGGG---GCTTACTCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA
SA17Lthesiodes	ACGTGATGGCCTCGTCGGGC-GGGG---GCTTACTCTCGTGGTC--GTGCC-TTCCCGCACAAACT-AACA
SA19Lcompton	ACGCGATGGCCTCGTCGAGC-GGGG---GCTTACTCTCGTGGTC--GTGCC-TTCCCGCACAAACA-AACA
SA1Lafricanum	ACGCGATGGCCTTGTGCGGC-GGGG---GCTTACTCTCGTGGTC--GTGCC-TTCCCGCGCAACT-AACA
SA3Lcuticarpum	ACGTGATGGCCTCGTCGGA-GGGG---GCTTACTCTCGTGGTC--ATGCC-TTCCCGCACAAACT-AACA
SA4Lgracile	ACGCGATGGCCTCGTCGGGC-GGGG---GCTTACTCTCGTGGTC--ATGCC-TTCCCGCACAAACT-AACA
SA5Lgracile	ACGCGATGGCCTCATCGGC-GGGG---GCTTACTCTCGTGGTC--GTGCC-TTCCCGCATAAACT-AACA
SA6Lgracile	ACGCGATGGCCTCGTCGGGC-GGGG---GCTTACTCTCGTGGTC--ATGCC-TTCCCGCATAAACT-AACA
SA7LinumDeHoop	ACGCGATGGCCTCATCGGC-GGGG---GCTTACTCTCGTGGTC--GTGCC-TTCCCGCATAAACT-AACA
SA8adustum	ACGCGATGGCCTTGTGCGGC-GGGG---GCTTACTCTCGTGGTC--ATGTC-TTCCCGCATAAACT-AACA
SA8Ladustum	ACGCGATGGCCTCATCGGC-GGGG---GCTTACTCTCGTGGTC--GTGCC-TTCCCGCATAAACT-AACA
SA9Lesterhuysena	ACGCGATGGCCTCCTCGGC-GGGG---GCTTACTCTCGTGGTC--GTGCC-TTCCCGCACAAACT-AACA
SDLinum	TCGCGATGGCCACGTCGTGTC-GGGG---GCTCGGCCCTTGTGGTC--GTTCC-TTCCCGCACAAACT-AACA

028Lberlandierif AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACGCCGTG--CTCGTGCCCGGTC-ACGGTGC  
032Lhudsonioides AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAGCGTCGCG--CTCGTGCCCGGTC-ACGGAGC  
034Lrupestre AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACGCCGTGTCTCGTGCCCGGTC-ACGGTGC  
035Lrigidum AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACGCCGTG--CTCGTGCCCGGTC-ACGGCGC  
036Lrupestre AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACGCCGTGTCTCGTGCCCGGTC-ACGGTGC  
037Lmexicanum AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTCGAGCGA-CGCCGTGTCTCGTGCCCGGTC-ACGGTGC  
038Llongipes AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGA-CGTGTGTCTCATGCCCCGGTC-ACGGTGC  
040Lcruciata AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGA-CGTGTGTCTCATGCCCCGGTC-ACGGTGC  
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055Laethiopicum AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGGGAGA-CACTGTGTCT-ATGCCCGGTC-ACGGTGC  
059Lcomptonii AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTCGAGCGA-CGCCGTGTCTCGTGCCCGGTC-ACGGTGC  
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066Ltenue AA-CCCCGGCGCGGA-ATGCGTCAAGGAATACT-TCGAGCGA-TGTCTTG-CCTCGTGCCCGGTC-TCGGTGC  
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149Lflavum AT-CCCCGACGCGGC-ATGCGTCAAGGAATACTTCGAGCGAACGTGCG-CCTTGTGCCCCGGTC-ACGGTGC  
151Lcatharticum AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTCGAGCGA-CGTGCGCTCTCGTGCCCGGTC-ACGGTGC  
156Ltenuifolium AACCCCGACGCGGC-ATGCGTCAAGGAATACTCCGAGCGA-CGTCTGTCTCATGCCCCGGTC-ACGGTGC  
157Ltauricum AT-CCCCGACGCGGC-ATGCGTCAAGGAATACTTCGAGCGAACGTGTC-CCTTGTGCCCCGGTC-ACGGTGC  
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164Larboeum AT-CCCCGACGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACGTGCG-CCTTGTGCCCCGGTC-TCGGTGC  
171Rlinioides AACCCCGACGCGGC-ATGCGTCAAGGAATACTATGAGCGA-CGACGTG-CTTCTGCCCCGGTC-ACGGTGC  
183Lpuberulum AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACGTGTC--CTCGTGCCCGGTC-ACGGTGC  
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246LrigidumKerr AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACGCCGTG--CTCGTGCCCGGTC-ACGGCGC  
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253Lelongatum	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAGCCCGGTG--CTCGTGCCCGGTC-ACGGTGC
256Lmedium	AA-CCCCGGCGCGGT-ATGCGTCAAGGAATACTTTGAGCGA-CGTCTGTCTCTGGTGCC-GGTC-ACGGTGC
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258Lquadrifolium	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTAGGGCGA-CGCCGTGTCTCTGTGCCCGGTC-ACGGTGC
266Lcarteri	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACCCGTG--CTCGTGCCCGGTC-ACGGCGC
268Larenicola	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACGCCGTGTCTCTGTGCCCGGTC-ACGGTGC
269Lwestii	AA-CCCCGGCGCGGT-ATGCGTCAAGGAATACTTTGATCGA-CGTTGTGTCTCTGGTGCC-GGTC-ACGGTGC
270Lorizabae	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTCGAGCAA-CGCCGTGTCTCTGTGCCCGGTC-ACGGTGC
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293Lguatemalense	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTCGAGCGA-CGCCGTGTCTCTGTGCCCGGTC-ACGGTGC
294Lcatharticum	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACATCGAGCGA-CGTCTCGCTCTCTGTGCCCGGTC-GCGGTGC
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310Llasiocarpum	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGA-CGTCTGTCTCTGTGCCCGGTC-ACGGTGC
401Lintercursum	AA-CCCCGGCGCGGT-ATGCGTCAAGGAATACTTTGAGCGA-CGTCTGTCTCTGTGGTGC-GGTC-ACGGTGC
AlRigidum	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACGCCGTG--CTCGTGCCCGGTC-ACGGTGC
F1Lbahamensecora	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACCCGTGTCTCTGTGCCCGGTC-ACGGTGC
F2Lcruciata	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTCGAGCGA-CGCCGTGTCTCTGTGCCCGGTC-CCGGTGC
F5Lnelsonii	AT-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTCGAGCGA-CGCCATGTCTCTGTGCCCGGTC-ACGGTGC
F6Ltenellum	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGA-CGTCTGTCTCTGTGCCCGATC-ACGGTGC
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L57Lflavum	AT-CCCCGACGCGGC-ATGCGTCAAGGAATACTTCGAGCGAACGCTCGC-CCTTGTGCCCGGTC-ACGGTGC
LinumSpYeso	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTAAGAGCGAACGCCGTG--CTCGTGCCCGGTC-ACGGTGC
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SA10Lesterhuysen	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTCGAGCGA-TGCCGTGTCTCTGTGCCCGGTC-ACGGTGC
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SA17Lthesiodes	AA-CCCCGACGCGGC-ATGCGTCAAGGAATACTTCGAGCGA-CGCCGTGTCTCTGTGCCCGGTC-ACGGTGC
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SA9Lesterhuysena	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTCGAGCGA-TGCCGTGTCTCTGTGCCCGGTC-ACGGTGC
SDLinum	AA-CCCCGGCGCGGC-ATGCGTCAAGGAATACTTTGAGCGAACCCGTG--CTCGTGCCCGGTC-ACGGTGC

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 042Lflagellare ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 047Hdrymarioides ACTGAGCGCGGCGATCCACCACATAA-TCACTACAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 048Hmicranthum ACTAAGCGCGGCGATCCACCAGTTAA-TCACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
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 137Lvolskensis TTTGGGCGTGGCGATCCACCAGTTAA-TTACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 139Lholstii TTTGGGCGTGGCGATCCACCAGTTAA-TTATTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
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 049Cselaginoides ATTGGGCGCGGCGATCCACCAGTTAA-CCACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 206Cselaginoides ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 210Lmacraei ATTAGGCGCGGCGATCCACCAGTTAA-TCACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 210Lmacraei ATTAGGCGCGGCGATCCACCAGTTAA-TCACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 215Lchamissonis ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 219Lscoparium ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 220Cselaginoides ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 221Lburkartii ATTGGGCGCGGCGATCCACCAGTTAA-TCAATATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 223Lkingii CTTGGGTGCGGCGATCCACCAGTTAA-TCACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 228Lneomexicanum ACTGGGCGCGGCGATCCACCAGTTAA-TCACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 231Lcompactum ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 232Laustrale ATTGGGCGCGGCGATCCACCAGTTAA-TGACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 233Laristatum ATCGGGCGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 234Lberlandieri ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 236Lrigidum ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 241Lberlandieri ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 242Lrupestre ATTGGGCGCGGCAATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 244Lhudsonioides ACTGAGCGCGGCGATCCACCAGTTAA-ACACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 246LrigidumKerr ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 249Limbricatum ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 250Lhudsonioides ACTGAGCGCGGCGATCCACCAGTTAA-ACACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 251Lrigidum ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 253Lelongatum ATCGGGCGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 256Lmedium ATTGGGCGCGGCGTCCATCACTTAA-TCAATATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 257Lmedium ATTGGGCGCGGCGTCCATCACTTAA-TCAATATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 258Lquadrifolium ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 266Lcarteri ATCGTGGCGGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 268Larenicola ATTGGGCGCGGCGATCCACCAGTTAA-TCACTATAATGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG  
 269Lwestii ATTGGGCGCGGCGTCCATCACTTAA-TCAATATAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCG



062Lheterostylum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

066Ltenue ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

067Lthunbergii ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

072Ltrigynum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

075Lstriatum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

083Sdigynum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

089Llittorale ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

091Loligophyllum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

123Lkingii ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

124Lfloridanum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

130Lvirginianum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

132Lsulcatum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

137Lvolkensii ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

139Lholstii ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

143Lorientale ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

149Lflavum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

151Lcatharticum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

156Ltenuifolium ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

157Ltauricum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

163Lleigans ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

164Larboreum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

171Lrinoides ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

183Lpuberulum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

184Lvernale ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

186Llundellii ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

187Lcremophyllu ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGWCCATCGAGTCTTTGAACGC

193Lbrevifolium ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

194Lprostratum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

195Lchamissonis ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

196Lerigeroides ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGWCCATCGAGTCTTTGAACGC

200Lcarneum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

201Lpringlei ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

049Cselaginoides ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

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210Lmacraei ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

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215Lchamissonis ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

219Lscoparium ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

220Cselaginoides ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

221Lburkartii ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

223Lkingii ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

228Lneomexicanum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

231Lcompactum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

232Laustale ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

233Laristatum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

234Lberlandieri ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

236Lrigidum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

241Lberlandieri ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

242Lrupestre ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

244Lhudsonioides ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

246LrigidumKerr ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

249Llimbricatum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

250Lhudsonioides ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

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253Lelongatum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

256Lmedium ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

257Lmedium ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

258Lquadrifolium ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

266Lcarteri ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

268Larenicola ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

269Lwestii ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

270Lorizabae ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

291Lrigidumfilif ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

293Lguatemalense ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

294Lcatharticum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

307Lalatum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

310Llasiocarpum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGC

401Lintercursum ATGAAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAAT-----



089Llittorale AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 091Loligophyllum AAGTTGCGCCYGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 123Lkingii AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 124Lfloridanum AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCTCA-TCCCCA  
 130Lvirginianum AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 132Lsulcatum AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 137Lvolskensis AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-ACCCCCA  
 139Lholstii AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-ACCCCCA  
 143Lorientale AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 149Lflavum AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 151Lcatharticum AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCTCCA  
 156Ltenuifolium AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-CCCCCA  
 157Ltauricum AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 163Lelegans AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 164Larboreum AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 171Lrinoides AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-ACCCCCA  
 183Lpuberulum AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 184Lvernale AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 186Llundellii AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
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 193Lbrevifolium AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
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 195Lchamissonis AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
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 200Lcarneum AAGTTGCGCCCCGAAGCCATCCGGCTAAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
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 049Cselaginoides AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
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 223Lkingii AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 228Lneomexicanum AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 231Lcompactum AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
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 233Laristatum AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 234Lberlandieri AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 236Lrigidum AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
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 242Lrueprestre AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 244Lhudsonioides AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 246LrigidumKerr AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
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 253Lelongatum AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 256Lmedium AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCTCG-TCCCCA  
 257Lmedium AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCTCG-TCCCCA  
 258Lquadrifolium AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA  
 266Lcarteri AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACTCAACGTCG-CCCCA-TCCCCA  
 268Larenicola AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 269Lwestii AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCTCA-TCCCCA  
 270Lorizabae AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 291Lrigidumfilif AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 293Lguatemalense AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 294Lcatharticum AAGTTGCGCCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCTCCA  
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 310Llasiocarpum AAGTTGCGCCCCGAAGCCATCCGGCTAAGGGCACGTCTGCCTGGGTGTACGCAACATCG--CCCA-TCCCCA  
 401Lintercursum -----GACTGAGGGCACGTCCGCTTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 AlLrigidum AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 FlLbahamensecora AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 F2Lcruciata AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA  
 F5Lnelsonii AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TTCCCCA  
 F6Ltenellum AAGTTGCGCCCCGAAGCCATCCGGCTAAGGGCACGTCTGCCTGGGTGTACGCAACGTCG--CCCA-TCCCCA  
 F7Lscabrellum AAGTTGCGCCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA



F8Ltenellum	AAGTTGCGCCCAAGCCATTCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCC-
L17Ltenuifolium	AAGTTGCGCCCGAAGCCATTCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-CCCCCA
L29Lmaritimum	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAATGTCG-CCCCA-TCCCCA
L43Lcatharticum	AAGTTGCGCCCGAAGCCATCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCTCCA
L57Lflavum	AAGTTGCGCCCGAAGCCATTCCGGTTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCC
LinumSpYeso	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA
PM1983Sdigynum	AAGTTGCGCCCGAAGCCACCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA
SA10Lesterhuysen	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA11LinumAgulhas	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA12Laethiopicum	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA13LinumspWindb	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA14Lvillosum	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGCGTACGCAACGTCG-CCACA-TCCCC
SA17Lthesiodes	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA19Lcompton	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA1Lafricanum	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGTAACGTCG-CCACA-TCCCCA
SA3Lcorticarpum	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA4Lgracile	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA5Lgracile	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA6Lgracile	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA7LinumDeHoop	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA8adustum	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA8Ladustum	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SA9Lesterhuysena	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCACA-TCCCCA
SDLinum	AAGTTGCGCCCGAAGCCATCCGGCTGAGGGCACGTCTGCCTGGGTGTACGCAACGTCG-CCCCA-TCCCCA

139Lholstii ACC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCGCTTGAGTCTTGC GGCTGGCCAAAA  
143Lorientale CTA--TG--GGA-GGGGAGCGGATGATGGCCTCCC-GGGG-GCTCTCTCTCGAGCCTCGCGGCTGGTCGAAA  
149Lflavum CTA--TG--GGA-GGGGAGCGGATGATGGCCTCCC-GGGG-GCTCTCTCTCGAGCCTCGCGGCTGGTCGAAA  
151Lcatharticum ATC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCACTCGAGCCTCGCGGTTGGCCGAAA  
156Ltenuifolium ACC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTAACTCGAGCCTCGCGGCTGGCCGAAA  
157Ltauricum CTA--TG--GGA-GGGGAGCGGATGATGGCCTCCC-GGGG-GCTCTCTCTCGAGCCTCGCGGCTGGTCGAAA  
163Lelegans CTA--TG--GGA-GGGGAGCGGATGATGGCCTCCC-GGGG-GCTCTCTCTCGAGCCTCGCGGTTGGTCGAAA  
164Larboreum CTA--TG--GGA-GGGGAGCGGATGATGGCCTCCC-GGGG-GCTCTCTCTCGAGCCTCGCGGCTGGTCGAAA  
171Lrinoides ATC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTC--TTGAGTCCCGCGGTTGGCCAAAA  
183Lpuberulum CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCATTCGAGCCTCGCGGCTGGCCGAAA  
184Lvernale CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
186Llundellii CCC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
187Lcremnophyllu TTC--G--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATGAGCCTTGGCGCTGGCCGAAA  
193Lbrevifolium CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTTGGCGCTGGTCGAAA  
194Lprostratum CTC--G--GGA--GGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATGAGCCTCGCGGCTGGCCGAAA  
195Lchamissonis CTC--GGGTGGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATGAGCCTCGCGGCTGGCCGAAA  
196Lerigeroides CTC--G--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATGAGCCTCGCGGCTGGCCGAAA  
200Lcarneum CTC--G--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATGAGCCTCGCGGCTGGCCGAAA  
201Lpringlei CTC--GG--GGA-GGGGAGCGGATGTTGGCTTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
049Cselaginoides CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
206Cselaginoides CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
210Lmacraei CTC--G--GGA--GGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATGAGCCTCGCGGCTGGCCAAAA  
210Lmacraei CTC--G--GGA--GGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATGAGCCTCGCGGCTGGCCAAAA  
215Lchamissonis CTC--GGGTGGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATGAGCCTCGCGGCTGGCCGAAA  
219Lscoparium CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTTGGCGCTGGCCAAAA  
220Cselaginoides CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
221Lburkartii CTC--GG--GGA-GGGGAGCGGATGTTGGCTTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
223Lkingii CCT--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCCCAATCGAGCCTCGCGGCTGGCCGAAA  
228Lneomexicanum CCC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCCCAATCGAGCCTCTCGGCTGGCCGAAA  
231Lcompactum CCC--GG--GGAGGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGTCCCGCGGCTGGCCGAAA  
232Laustale CCC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
233Laristatum CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
234Lberlandieri CYC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAACCGAGYCYCGCGCTGGCCGAAA  
236Lrigidum CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAACCGAGTCTCGCGGCTGGCCGAAA  
241Lberlandieri CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAACCGAGTCTCGCGGCTGGCCGAAA  
242Lrupestre CTT--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
244Lhudsonioides CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCACGATCGAGCCTCGCGGCTGGCCGAAA  
246LrigidumKerr CCC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCCCGCGGCTGGCCGAAA  
249Limbricatum CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
250Lhudsonioides CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCCA--CGAGCCTCGCGGCTGGCCGAAA  
251Lrigidum CCC--GG--GGAGGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGTCCCGCGGCTGGCCGAAA  
253Lelongatum CCC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
256Lmedium CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCTC-GTGG-GCTCCTAATCGAGCCTTGGCGTTGGCCGAAA  
257Lmedium CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCTC-GTGG-GCTCCTAATCGAGCCTTGGCGTTGGCCRAAA  
258Lquadrifolium CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
266Lcarteri CCC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
268Larenicola CTT--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTTGGCGTTGGCCGAAA  
269Lwestii CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCTC-GTGG-GCTCCTAATCGAGCTTGGCGATTGGCCGAAA  
270Lorizabae CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCTC-GGGG-GCTCTCAATGAGCCTTGGCGCTGGCCGAAA  
291Lrigidumfilif CAC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCCCGCGGCTGGCCGAAA  
293Lguatemalense CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCTC-GGGG-GCTCTCAATGAGCCTTGGCGCTGGCCGAAA  
294Lcatharticum ATC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCACTCGAGCCTCGCGGTTGGCCGAAA  
307Lalatum CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
310Llasiocarpum CTC--AG--GGA--GGGAGCGGATGTTGGCCTCCCCTGGCGCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
401Lintercursum CTT--GG--GGA-GGGGAGCGGATGTTGGCCTCTC-GTGG-GCTCCTAATCGAGCCTTGCAGTTGGCCGAAA  
AlLrigidum CCC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
FlLbahamensecora CTT--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
F2Lcruciata CTC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTGTCAATCAAGCCTCACGTTGGCCGAAA  
F5Lnelsonii CTT--GG--GGA-GGGGGGCGGATATTGGCCTCCC-GGGG-GCTCTTAATCAAGCCTCGCGGTTGGCCAAAA  
F6Ltenellum CTC--AG--GGA--GGGAGCGGATGTTGGCCTCCCCTGGCGCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
F7Lscabrellum CTT--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCAATCGAGCCTCGCGGCTGGCCGAAA  
F8Ltenellum CTT--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GAGTAGCTCAATGAGCCTCACGTTGGCCGAAA  
L17Ltenuifolium ACC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTAACTCGAGCCTCGCGGCTGGCCGAAA  
L29Lmaritimum CTC--GG--GGA-TAGGAGCGGATGTTGGCCTCCC-GGGG-GCTC---TTAGCCTTGGCGTTGGCCGAAA  
L43Lcatharticum ATC--GG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCACTCGAGCCTCGCGGTTGGCCGAAA  
L57Lflavum CTA--TG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCTCGAGCCTCGCGGCTGGTCGAAA  
LinumSpYeso CTC--GG--GAA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCTCTATCGAGCCTCGCGGCTGGTCGAAA  
PM1983Sdigynum CCG--TG--GGA-GGGGAGCGGATGTTGGCCTCCC-GGGG-GCTCCCAACCGAGCCCCGCGGCTGGCCGAAA



164Larboreum A-GAAAGTCGTCGGCCTAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
171Rlinoides A-GAAAGTCGTCGGCAGAGGATGCCGTAGCGAGCTGTGGTTGAAATTTCACT--AGTCTCATTTGCTACGTG  
183Lpuberulum A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
184Lvernale A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
186Llundellii A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
187Lcremophyllu A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
193Lbrevifolium A-AGAATTCGTCGGCAGAGGATGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGTTGGCTGCGTG  
194Lprostratum A-AAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
195Lchamissonis A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
196Lerigeroides A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
200Lcarneum A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGG-----  
201Lpringlei A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGATGGCTGCGTG  
049Cselaginoides A-GATATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
206Cselaginoides A-GATATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
210Lmacraei A--AAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
210Lmacraei A--AAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACA-----  
215Lchamissonis A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
219Lscoparium A-AGAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGTTAGCTGCGTG  
220Cselaginoides A-GATATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
221Lburkartii A-GATTTTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGATGGCTGCGTG  
223Lkingii A-GAAATTCATCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTCAGCTGCGTG  
228Lneomexicanum A-GAAATTCATCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTCGGCTGCGTG  
231Lcompactum A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
232Liberale A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
233Laristatum A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
234Lberlandieri A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
236Lrigidum A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
241Lberlandieri A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
242Lrupestre A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGACGGCTGCGTG  
244Lhudsonioides A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
246LrigidumKerr A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
249Limbricatum A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGATGGCTGCGTG  
250Lhudsonioides A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
251Lrigidum A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
253Lelongatum A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
256Lmedium T-GAAATTCGTCGGCATAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCATGGCTGCGTG  
257Lmedium T-GAAATTCGTCGGCATAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCATGGCTGCGTG  
258Lquadrifolium A-GAAAGTCATCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTCGGCTGCGTG  
266Lcarteri A-GGAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
268Larenicola A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGACGGCTGCGTG  
269Lwestii T-GAAATTCGTTGGCATAGGATGCCGTTGCGAGCTGTGGTTGAAATTAAACTGGAGTCTCATGGCTGCGTG  
270Lorizabae A-GAAAGTCGTTGGCAGAGGACGCCGTTGCGAGCTATGGTTGAAATTACACTAGAGCGCTCGTTGGCTGCGTG  
291Lrigidumfilif A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
293Lguatemalense A-GAAAGTCGTTGGCAGAGGACGCCGTTGCGAGCTATGGTTGAAATTACACTAGAGCCTCGTTGGCTGCGTG  
294Lcatharticum AGGAAAGTCGTCGGCTGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
307Lalatum A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
310Llasiocarpum A-GAAATTCATCGGTAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
401Lintercursum T-TAAATTCGTCGGCATAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCATGGCTGCGTG  
AlLrigidum A-GATATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
FlLbahamensecora A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGACGGCTGCGTG  
F2Lcruciata A-CAATTCGTCGTCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCAGCGTG  
F5Lnelsonii A-GAATTCGTCGTCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
F6Ltenellum A-GAAATTCATCGGTAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
F7Lscabrellum A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGATGGCTGCGTG  
F8Ltenellum A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTATGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
L17Ltenuifolium A-GAAAGTCGTCGGCTGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
L29Lmaritimum A-GAAAGTCGTCGGCTGAGSACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTCGCATACGTG  
L43Lcatharticum AGGAAAGTCGTCGGCTGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
L57Lflavum A-GAAAGTCGTCGGCTAGAACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
LinumSpYeso A-GAAATTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTGGAGTCTCGATGGCTGCGTG  
PM1983Sdigynum A-GAAAGTCGTCGGCAGGGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTCAGCTGCGTG  
SA10Lesterhuysen A-GAAAGTCATCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
SA11LinumAgulhas A-GAAAGTCATCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
SA12Laethiopicum A-GAAAGTCATCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
SA13LinumspWindb A-GAAAGTCATCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
SA14Lvillosum A-GAAAGTCGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
SA17Lthesiodes A-GAAAGTCATCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG  
SA19Lcompton A-GAAAGTTGTCGGCAGAGGACGCCGTTGCGAGCTGTGGTTGAAATTACACTAGAGTCTCGTTGGCTGCGTG



194Lprostratum CGACCTCCTCCGGCGGCACCAC-GATGGACCTCGAGGGAC--A-TT-----TGAACCTTCAAAAATGC  
195Lchamissonis CGATCCTCCGTCCGGGGCACCAC-GATGGACC-----TGAACCTTCAAAAATGC  
196Lerigeroides CGATCCTCCGTAGGCGGCAACAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTAAAAAATGC  
200Lcarneum -----TGAACCTTCAAAAATGC  
201Lpringlei TGATCCTCCGTCCGGTGGCACCAC-TAAGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
049Cselaginoides CGATCCTCCGTTCGATGGCACCAC-GATGGACCTTGAAGGGAC--A-GT-----TGAACCTTCAAAAATGC  
206Cselaginoides CGATCCTCCGTTCGATGGCACCAC-GATGGACCTTGAAGGGAC--A-GT-----TGAACCTTCAAAAATGC  
210Lmacraei CGATCCTTTGTCCGGCGGCACCAT-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
210Lmacraei -----TGAACCTTCAAAAATGC  
215Lchamissonis CGATCCTCCGTCCGGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
219Lscoparium CGATCCTCCGTTCGATGGGACAAAC-TATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
220Cselaginoides CGATCCTCCGTTCGATGGCACCAC-GATGGACCTTGAAGGGAC--A-GT-----TGAACCTTCAAAAATGC  
221Lburkartii CGATCCTCCGTTCGATGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
223Lkingii CGATCCTCCGTTCGATGGCACCAC-TATGGACCTTACGGGAC--A-AC-----CGATCCTTAAAAA-GC  
228Lneomexicanum CGATCCTCCGCCGATGGCACCAT-TATGGACCTTTCGGGAC--A-AT-----TGATCCTT-AAAAATGC  
231Lcompactum CGATCCTCCGTTCGATGGTACCCC-AAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
232Laustrale CGATCCTCCGTTCGATGTTACCCC-CAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
233Laristatum CGATCCTCCGTTCGATGGTACCCC-AAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
234Lberlandieri CGATCCTCCGTTCGATGGTACCCC-CAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
236Lrigidum CGATCCTCCGTTCGATGGTACCCC-CAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
241Lberlandieri CGATCCTCCGTTCGATGGTACCCC-CAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
242Lrupestre CGATCCTCCGTTCGACGGCACCAC-TAAGGACCTCGAGGGAC--A-GT-----TGAACCTTCAAAAATGC  
244Lhudsonioides CGATCCTCCGTTCGACGGTACCCC-TGAGGACCTCGAGGGAC--A-AG-----TGAGCCTTCAAAAATGC  
246LrigidumKerr CGATCCTCCGCCGATGGTACCCC-AAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
249Limbricatum CGATCCTCCGTTCGATGGTACTCC-TAAGGACCTCGAGGGAC--A-AT-----TGAGCCTTCAAAAATGC  
250Lhudsonioides CGATCCTCCGTTCGACGGTACCCC-TGAGGACCTCGAGGGAC--A-AG-----TGAGCCTTCAAAAATGC  
251Lrigidum CGATCCTCCGTTCGATGGTACCCC-AAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
253Lelongatum CGATCCTCCGTTCGATGGTACCCC-AAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
256Lmedium CGATCCTTTGTTCGATGTGACCAA-GATGGACCTCGAGGGAC--AAAT-----TGAGCCTTCAAAAGATGC  
257Lmedium CGATCCTTTGTTCGATGTGACCAA-GATGGACCTCGAGGGAC--AA-AT-----TGAGCCTTCAAAAGATGC  
258Lquadrifolium CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
266Lcarteri CGATCCTCCGTTCGATGGTACCCA-CAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
268Larenicola CGATCCTCCGTTCGACGGCACCAC-TAAGGACCTCGAGGGAC--A-GT-----TGAACCTTCAAAAATGC  
269Lwestii CGATCCTTTGCCGATGTGCCAC-GATGGACCTCGAGGGAC--A-AT-----TGAGCCTTCAAAAATGC  
270Lorizabae CGATCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
291Lrigidumfilif CGATCCTCCGTTCGATGGTACCCC-AAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
293Lguatemalense CGATCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
294Lcatharticum CGCTCCTTTGTTCGATGACACCAT-GATGGACCTCGAGGGACCAATAA-----CGGACCTTCAAAAATGC  
307Llalatium CGATCCTCCGTTCGATGGTACCCC-AAAGGACCACGAGGGAC--A-GC-----TGAACCTTCAAAAATGC  
310Llasiocarpum CGATCCTTCGTTGATGGAACCAT-GATGGACCTC-----TGAACCTTCAAAAATGC  
401Lintercursum CGATCCTTTGTTCGATGTGACCAA-GATGGACCTCGAGGGAC--AA-AT-----TGAGCCTTCAAAAATGC  
AlLrigidum CGATCCTCCGTTCGATGGTACCCC-AAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
F1Lbahamensecora CGATCCTCCGTTCGACGGCACCAC-TAAGGACCTCGAGGGAC--A-GT-----TGAACCTTCAAAAATGC  
F2Lcruciata CGATCCTTTGTTCGATGGCCCCAT-GATGGACCTCGAGGGAC--AA-----TTGACCTTCAAAAATGC  
F5Lnelsonii CGATCCTTCGTTGATGGCCTCAT-GATGGACCTCGAGGGAC--AA-----TTGACCTTCAAAAATGC  
F6Ltenellum CGATCCTTCGTTGATGGAACCAT-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
F7Lscabrellum CGATCCTCCGTTCGACGGCACCAC-TAAGGACCTCGAGGGAC--A-GT-----TGAACCTTCAAAAATGC  
F8Ltenellum CGATCCTTTGTTCGATGGAACCAT-GATGGACCTCGAGGGAC--A-AT-----TTATCCTTCAAAAATGC  
L17Llenuifolium CGCTCCTCCGCCGAGGACACCAC-GATGGACCTCGAGGGACATA-AA-----TGGACCTTCAAAAATGC  
L29Lmaritimum TGCTCCTTGGTTGATGTCTACAC-AATGGACCTGAGGGACTAA-----TGTACCTTCAAAAATGC  
L43Lcatharticum CGCTCCTTTGTTCGATGACACCAT-GATGGACCTCGAGGGACCAATAA-----CGGACCTTCAAAAATGC  
L57Lflavum CGCTCCTGGGTTCGAGGACRCCAT-GATGGACCTCGAGGGAC--G-AA-----TGGACCTTCAAAAATGC  
LinumSpYeso CGATCCTCCGTTCGATGGTACCCC-CAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC  
PM1983Sdigynum CGCTCCTTTGTTCGAGGACGCCAC-GACGGACCTCGAGGGAC--A-AC-----CGAGCCTTCAAAAATGC  
SA10Lesterhuysen CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA11LinumAgulhas CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA12Laethiopicum CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA13LinumspWindb CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA14Lvillosum CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA17Lthesioides CGCTCCTCCGTTCGAGGGCACCAT-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA19Lcompton CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA1Laficanum CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA3Lacuticarpum CGCTCCTTCGTCGAGGGCACCAC-GATGGACCTCAAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA4Lgracile CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA5Lgracile CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA6Lgracile CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA7LinumDeHoop CGCTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC  
SA8adustum CGTTCCTCCGTTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTCAAAAATGC

SA8Ladustum	CGCTCCTCCGTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTAAAAAATGC
SA9Lesterhuysena	CGCTCCTCCGTCGAGGGCACCAC-GATGGACCTCGAGGGAC--A-AT-----TGAACCTTTACAAATGC
SDLinum	CGATCCTCCGTCGATGGTACCCC-AAAGGACCACGAGGGAC--A-AC-----TGAACCTTCAAAAATGC

210Lmacraei	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
210Lmacraei	-----
215Lchamissonis	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCA-----
219Lscoparium	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATA-----
220Cselaginoides	GA-CCCCAGGTCAGGCGGGAAGCACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
221Lburkartii	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGG-
223Lkingii	GA-CCCCAGGTCAGGCG-----
228Lneomexicanum	GA-CCCCAGGTCAGGCGG-----
231Lcompactum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
232Laustrale	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
233Laristatum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
234Lberlandieri	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAAT-----
236Lrigidum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
241Lberlandieri	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
242Lrupestre	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATA-----
244Lhudsonioides	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
246LrigidumKerr	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGC-----
249Llimbricatum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
250Lhudsonioides	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
251Lrigidum	GA-CCCCAGGT-----
253Lelongatum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
256Lmedium	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
257Lmedium	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAA-----
258Lquadrifolium	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAA-----
266Lcarteri	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
268Larenicola	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
269Lwestii	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
270Lorizabae	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
291Lrigidumfilif	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCA-----
293Lguatemalense	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
294Lcatharticum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTA-----
307Llalatium	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
310Llasiocarpum	-----
401Lintercursum	GA-CCCCAGGTCAGGCGGGAACACCCACTGAGTTTAAGCATATCAATAAGCGGAGGA
AlLrigidum	GG-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
F1Lbahamensecora	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAA-----
F2Lcruciata	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
F5Lnelsonii	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
F6Ltenellum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAG-----
F7Lscabrellum	-----
F8Ltenellum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAA-----
L17Ltenuifolium	-----
L29Lmaritimum	-----
L43Lcatharticum	-----
L57Lflavum	-----
LinumSpYeso	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
PM1983Sdigynum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA10Lesterhuysen	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAA-----
SA11LinumAgulhas	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA12Laethiopicum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA13LinumspWindb	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATAT-----
SA14Lvillosum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA17Lthesiodes	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA19Lcompton	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA1Lafricanum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA3Lacuticarpum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA4Lgracile	GA-CCCCAGGTCAGGCG-----
SA5Lgracile	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA6Lgracile	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA7LinumDeHoop	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA8adustum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA8Ladustum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA
SA9Lesterhuysena	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGG-
SDLinum	GA-CCCCAGGTCAGGCGGGAACACCCGCTGAGTTTAAGCATATCAATAAGCGGAGGA



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## Vita

Joshua Robert McDill was born in Vallejo, California, but was raised in the town of Napa. Following an unremarkable primary education, he moved to San Francisco, where, as an unremarkable undergraduate at San Francisco State University, he was introduced to the study of botany by several charismatic professors who made it all seem fun. Following a botany internship at the California Academy of Sciences, he entered graduate school at S.F.S.U. under the supervision of Dr. Robert Patterson, and received a M.A. in Ecology and Systematic Biology in 2001 with a thesis entitled “Evolutionary Trends in the Leaf and Stem Anatomy of *Linanthus* and Related Genera (Polemoniaceae).” While collecting *Linanthus* throughout California, he became familiar with another genus, *Hesperolinon*, which initiated a fateful interest in the Linaceae. At S.F.S.U., he served as a teaching assistant for courses including The World of Plants, Electron Microscopy, and The Sierra Nevada Flora. He entered the Graduate School of the University of Texas at Austin in the Fall of 2001, in the Plant Biology Graduate Program. While at U.T., he was a teaching assistant for courses including Plant Anatomy, Survey of the Plant Kingdom, and Native Plants, and served as a graduate research assistant for Dr. Beryl Simpson. He received a University Pre-emptive Fellowship for the 2001-2002 academic year, a National Science Foundation Dissertation Improvement Grant in 2005, and a University Continuing Fellowship for the 2006-2007 academic year. He resides in Austin, TX, with his husband, Marc Caddell.

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